73732

From:

Hashemi, Shar

Sent: To:

Monday, August 19, 2002 11:21 PM

STIC-Biotech/ChemLib

Subject:

Please search ALL NA including Pending SEQ. ID NO: 9, 29-31, 33 (five sequences) for

application 10/077,383

Please search ALL NA including Pending SEQ. ID NO: 9, 29-31, 33 (five sequences) for application 10/077,383

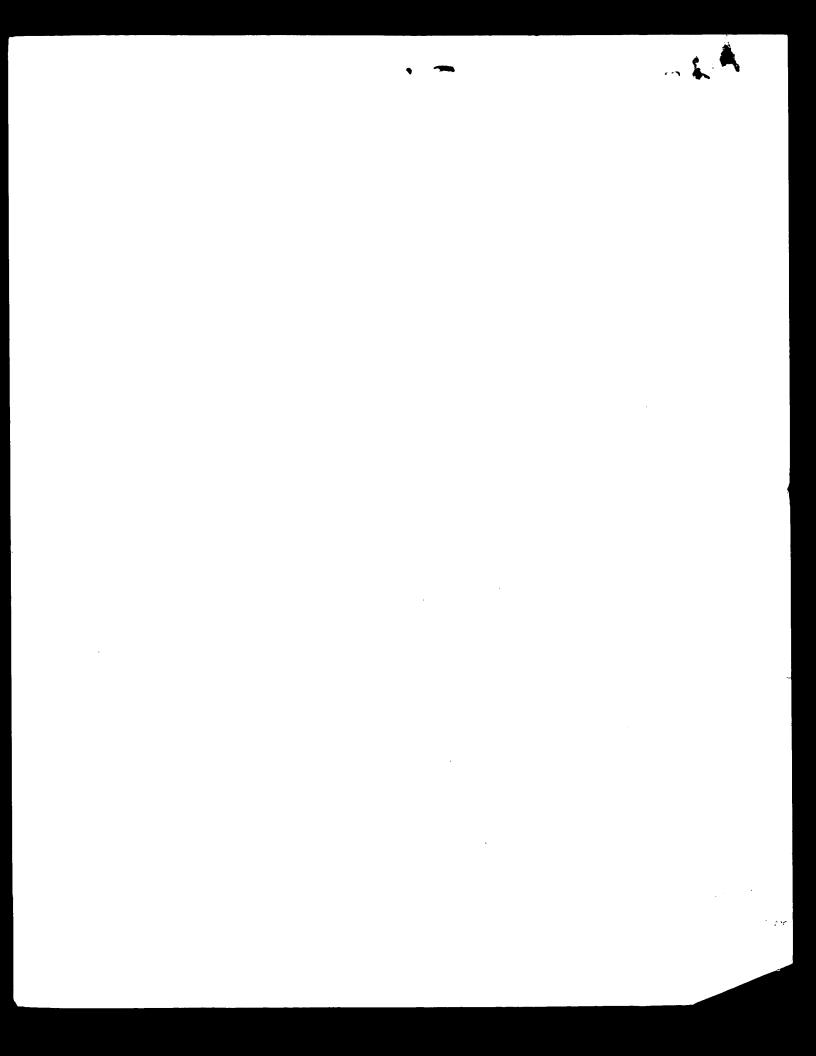
Thanks. Examiner Hashemi CM1 12D04 703-305-4840

> The segs. disclosed for Seg. IDs 9,30\$ 33 consists only of "N"s; therefore, no query match > 0 was jocated/retrieved. Beverly 1E05 C8-4994

Searcher:
Phone:
Location:
Date Picked Up:
Date Completed:
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

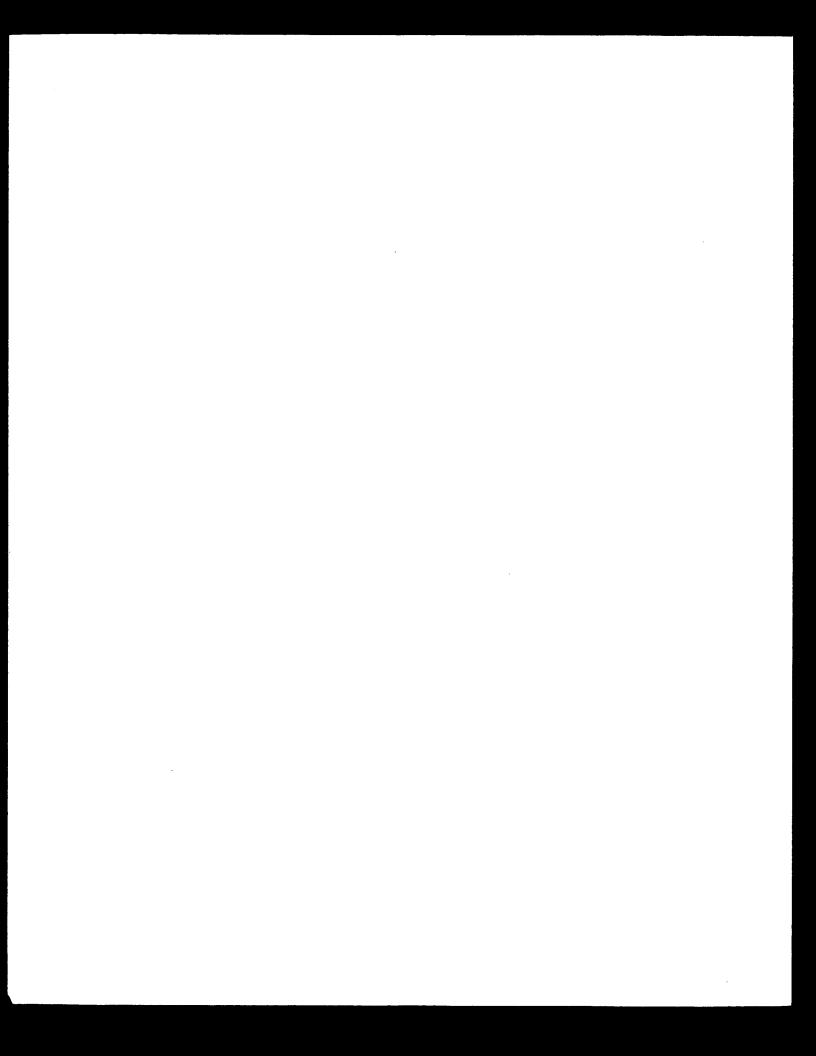
VENDOR/COST (where applic.)
STN:	_
DIALOG:	_
Questel/Orbit:	_
DRLink:	_
Lexis/Nexis:	
Sequence Sys.:	_
WWW/Internet:	_
Other (specify):	_



SEARCH REQUEST FORM

Requestor's Name:			Serial Number:			
Date:		Phone:		_ Art Unit:		
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Search Topic: Please write a detaile terms that may have a please attach a copy of	special meaning, G	ive examples or relev	ent citations, autho	rs, keywords, etc.,	if known. For sequ	ine any ences,
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Number of Searches:		·	A.A. Seque	nce	SDC	
Number of Databases	:	 .	Structure Bibliograph	.	DARC/Qu Other	

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'cgg1_6'ptodata/2'ppia/10509_COMB.seq:*
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1: /cgy12_6/ptodata/Z/pna_VISCOMB.seq:*

3: /cgn2_6/ptodata/Z/pna_VISO6_COMB.seq:*

4: /cgn2_6/ptodata/Z/pna_VISO7_COMB.seq:*

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5: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

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8: /cgn2_6/ptodata/Z/pna_VISO81_COMB.seq:*

9: /cgn2_6/ptodata/Z/pna_VISO81_COMB.seq:*

10: /cgn2_6/ptodata/Z/pna_VISO81_COMB.seq:*

11: /cgn2_6/ptodata/Z/pna_VISO81_COMB.seq:*

11: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

11: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

12: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

13: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

14: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

15: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

17: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*

17: /cgn2_6/ptodata/Z/pna_VISO80_COMB.seq:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                    OM nucleic - nucleic search, using sw model.
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Listing first 45 summaries
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Sequence:
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/ cgn2_6/ptodata/2/pma/US6006_C0M8_Seq:
/ cgn2_6/ptodata/2/pma/US6006_C0M8_Seq:
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/ cgn2_6/ptodata/2/pma/US6009_C0M8_Seq:
/ cgn2_6/ptodata/2/pma/US6010_C0M8_Seq:
/ cgn2_6/ptodata/2/pma/US6011_C0M8_Seq:
/ cgn2_6/ptodata/2/pma/US6012_C0M8_Seq:
/ cgn2_6/ptodata/2/pma/US6012_C0M8_Seq:
/ cgn2_6/ptodata/2/pma/US6012_C0M8_Seq:
/ cgn2_6/ptodata/2/pma/US6022_C0M8_Seq:
/cgn2_6/ptodata/2/pna/US6005_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	DI	-0893	893-00977	-068	-08	-08-154	-08-154-7	-08-154	-08-154-792A-	-08-163-4	8-163-	-08-163-449A-	-08-163-449A-	US-08-208-572-10		US-08-237-491-6868	-237-491-68	-296-1	296-757B-7	-177-	09-177-650-	-306-333A-1	-09-306-333A-	09-625-102-3	-09-625-1	-09-625-	-09-625-102-4	-09-625-102-	-09-625-102-58	-09-625-102-60	-09-625-1	US-09-625-102-614
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/cgn2_6/ptodata/2/pna/US101_COMB.seq:* /cgn2_6/ptodata/2/pna/US6000_COMB.seq:

/cgn2_6/ptodata/2/pna/US6001_COMB.seq:*/cgn2_6/ptodata/2/pna/US6002_COMB.seq:*/cgn2_6/ptodata/2/pna/US6003_COMB.seq:*/cgn2_6/ptodata/2/pna/US6004_COMB.seq:*/cgn2_6/ptodata/2/pna/US6004_COMB.seq:*/

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TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
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APPLICATION NUMBER: 07/659,980
                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                      Floppy disk
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TYPE: NUCLEIC ACID
STRANDEDNESS: double
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MEDIUM TYPE: Floppy C
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                                                                             Newport Beach
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CLASSIFICATION: 435
                                                                                                                                                                                                                                    APPLICATION NUMBER:
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STATE: CA
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                                                                                                            USA
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PCT-US93-00977-702
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                                                                                                            COUNTRY:
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                                                            Sequence 895, App
Sequence 982, App
Sequence 1048, App
Sequence 1048, App
Sequence 1352, App
Sequence 1352, App
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Sequence 51773, A
Sequence 51841, A
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; GENRRAL INFORMATION:
    TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
    NUMBER OF SEQUENCES: 711
    CORRESPONDENCE ADDRESS:
     Sequence
                  Sequence
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                                                           US-09-625-102-895
US-09-625-102-982
US-09-625-102-982
US-09-625-102-1048
US-09-625-102-1352
US-09-625-102-1352
US-09-634-3068-51773
US-09-634-3068-51773
                                                                                                                                                                                                                                                                                                                                                                                                          SEE: Knobbe, Martens, Olson, and Bear
: 620 Newport Center Dr. Sixteenth Floor
Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                       US-09-634-306B-51841
                                 US-09-625-102-836
   US-09-625-102-614
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-9404
TELEFAX: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 702: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: CDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 34
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Matches 0; Conservative
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STRANDEDNESS: double
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PCT-US93-00977-702/c
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PCT-US93-00977-702
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GENERAL INFORMATION:
APPLICANT: MUENCHAU, DARYL, ET AL.
TITLE OF INVENTION: METHOD FOR GENERATING APTAMERS OF
TITLE OF INVENTION: NON-PREDETERMINED OR SUBSTANTIALLY
TITLE OF INVENTION: NON-PREDETERMINED SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.0%; Score 0; DB 1; Length 1; 0.0%; Pred. No. 0; tive 0; Mismatches 1; Indels
NUMBER OF SEQUENCES: 711
CORRESPONDENCE ADDRESS:
CONDESSEE: Knobbe, Martens, Olson, and Bear
STREET: 620 Newport Center Dr. Sixteenth Floor
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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19930129
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                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Altman, Daniel E. REGISTRATION NUMBER: 34,115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: HIT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 714-760-9502
INFORMATION FOR SEQ ID NO: 702:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: LYON & LYON
STREET: 611 West Sixth St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: CDNA to mRNA
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us-10-077-383-9.rnpm

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0; Gaps
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GENERAL INFORMATION:

TITLE OF INVENTION: METHOD FOR GENERATING APTAMERS OF

TITLE OF INVENTION: MCHOD FOR GENERATING APTAMERS OF

TITLE OF INVENTION: MON-PREDETERMINED OR SUBSTANTIALLY

TITLE OF INVENTION: NON-PREDETERMINED SEQUENCES

NUMBER OF SEQUENCES: 26

CORRESPONDENCE ADDRESS:

ADDRESSEE: LYON & LYON

STREET: 611 West Sixth St.
                                                                                                                                                                                                                                                                                                                                                                         COTHER INFORMATION: where N also may be isoguanine or OCHER INFORMATION: isocytostine US-08-068-298-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.0%; Score 0; DB 4; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/068,298
                                                                                            193/051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         193/051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/659,980
FILING DATE: February 21, 1991
ATTORNEY/AGENT INFORMATION:
FILING DATE: February 21, 1991 ATTORNEY/AGENT INFORMATION:
                                        NAME: CONSALVI, MARY S
REGISTRATION NUMBER: 32,13,
REFERENCE/DOCKET NUMBER: 193,0
TELECOMMUNICATION INFORMATION:
TELEPRING: 213-499,1600
TELEFAX: 213-95-0440
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: CONSALVI, MARY S

REGISTRATION NUMBER: 32,212

REFERRACE/POCKET NUMBER: 193/O

TELECOMMUNICATION INFORMATION:

TELEPHONE: 213-499-1600

TELEFAX: 213-955-0440
                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 213-955-0440 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                   : 1 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                       single
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Los Angeles
                                                                                                                                                                                                                                                                                                           linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-068-298-17/c
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0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF INVENTION: and Uses Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
                                                                                                    Query Match 0.0%; Score 0; DB 4; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
OTHER INFORMATION: where N also may be isoguanine or OTHER INFORMATION: isocytostine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/154,792A FILING DATE: 17-NOV-1993 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                    ; Sequence 9, Application US/08154792A ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          702-08-154-792A-9/c

72-08-164-792A-9/c

56-08-164-792A-9/c

56-08-164-792A-9/c

GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 4482.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: (617) 523-6440
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kolodner, Richard D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34,235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-154-792A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
REGISTRATION NUMBER: 34,2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Boston
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    ; OTHER INFC
; OTHER INFC
US-08-068-298-17
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US-08-154-792A-24
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GENERAL INFORMATION:
APPLICANT: Recland Nobert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Gene Products,
TITLE OF INVENTION: and Uses Therefor
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF INVENTION: and Uses Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSE: David 6. Conlin; DIKE, BRONSTEIN, ROBERTS &
ADDRESSEE: CUSHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            / Match 0.0%; Score 0; DB 5; Length 1; Local Similarity 0.0%; Pred. No. 0; nes 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                       FILING DATE: US/08/154,792A CLASTAING DATE: 17-NOV-1993
CLASTAINO: 435
TORNEY FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 34,235
REGISTRATION NUMBER: 44824
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-6440
TELEFAX: (617) 523-6440
TELEFAX: (617) 523-6440
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-154-792A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          130 Water Street
                                                                                                                                                        130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Resnick, David S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                 NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                           Boston
                                                                                                                                                                                                                   USA
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US-08-154-792A-24
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                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
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                                                                                                                                                                                                STATE:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS &
ADDRESSEE: CUSHWAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 0; DB 5; Length 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PR-POS-MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.0%; Score 0; DB 5; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/154,792A FILING DATE: 17-NOV-1993
APPLICATION NUMBER: US/08/154,792A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24, Application US/08154792A GENERAL INFORMATION:
                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEPA: (617) 523-6440
TELET: 200291 STRE UR
INFORMATION POR SEX ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: I base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kolodner, Richard D. APPLICANT: Reenan, Robert A.G. APPLICANT: Fishel, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 44824
TELECOMMUNICATION INFORMATION:
TELECHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: (617) 523-6440
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                               ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
REGISTRATION NUMBER: 34,235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34,235
                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-154-792A-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S
REGISTRATION NUMBER: 34,2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  130 Water Street
                   FILING DATE: 17-NOV-1993
CLASSIFICATION: 435
                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                               nucleic acid
EDNESS: single
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nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                linear
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                              NAME: Resentck, David S.
REGISTRATION NUMBER: 34,234
REGISTRATION NUMBER: 44824
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACIERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 130 Water Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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CLASSIFICATION:
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                                                                                                                           Boston
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US-08-163-449A-24
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                                                                                                                                                                 COUNTRY:
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                                          1; Indels
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Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                            Sequence 9, Application US/08163449A
GENERAL INFORMATION:
APPLICANT: Recenn, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES THEREFOR NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: 07-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: DATID G. CONLIN; DIKE, ERONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           44824-CIP-2
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GENERAL INFORMATION:
APPLICANT: Kednar, Richard D.
APPLICANT: Fishel, Richard
APPLICANT: Fishel, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 34,234
REFERENCE/DOCKET NUMBER: 44824
TELECOMMUNICATION INFORMATION:
TELEPRIONE: (617) 523-5400
TELERAX: (617) 523-6440
TELERAX: (617) 523-6440
TELERAX: 06291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Resnick, David S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-163-449A-9/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24, Application US/08163449A
GENERAL INFORMATION:
APPLICANT: RCOLONC:
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TILE OF INVENTION:
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AUSBATCH
INVENTION: AUSBATCH
NUMBER OF SOURNCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PAPLICATION NUMBER: US/08/163,449A
FILING DATE: 07-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
COFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES THEREFOR
COMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
                                                                                                                                            ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN
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ER: 44824-CIP-2
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Gaps

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0; Gaps
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APPLICANT: Balcewicz-Sablinska M.D., Maria K.
APPLICANT: Balcewicz-Sablinska M.D., Maria K.
APPLICANT: Newman Ph.D., Gale W.
ATPLE OF INVENTION: CONTROL OF HIV GROWTH WITH ECEF AND TITLE OF INVENTION: ECEF
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESSE: CHOATE, & STEWART
STREET: EXCHANGE Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10, Application US/08208572
GENERAL INFORMATION:
APPLICANT: Silberstein Ph.D., David S.
APPLICANT: Balcawicz-Sablinska M.D., Maria K.
APPLICANT: Newman Ph.D., Gale W.
TITLE OF INVENTION: CONTROL OF HIV GROWTH WITH ECEF AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
                                                                                                           1; Indels
                                                          Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: 092662-002
TELECOMONICIONION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/208,572
                                                                                                                                                                                                                                                                                                                Sequence 10, Application US/08208572 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kaplan Esq., Warren A.
REGISTRATION NUMBER: 34,199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPAN. (617) 227-5020
TELEPAN. (617) 227-7566
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , MOLECULE TYPE: DNA (genomic) US-08-208-572-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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US-08-163-449A-24
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CURRENT APPLICATION DATA:
PAPLICATION NUMBER: US/08/163,449A
FILING DATE: 07-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Recenan, Robert A.G.
APPLICANT: Recenan, Robert B.G.
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES THEREFOR
NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                 REFERENCE/DOCKET NUMBER: 44824-CIP-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFX: 202915 STRE UR
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        44824-CIP-2
APPLICATION NUMBER: US/08/163,449A FILING DATE: 07-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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TELECHONE: (617) 523-3400
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                   NAME: Resnick, David S. REGISTRATION NUMBER: 34,234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34,234
                                                                                                                                                                                                                                                                                                                                                                                    ) MOLECULE TYPE: DNA (genomic) US-08-163-449A-24
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                                                               ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                QUENCE CHANNEL LENGTH: 1 base pairs TYPE: nucleic acid cannandEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
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                                            CLASSIFICATION:
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APPLICANT: Sellhamer, Angelo
APPLICANT: Delegeane, Angelo
APPLICANT: Jernigan, Colleen
APPLICANT: Levine, Wendy B.
TITLE OF INVENTION: SHEAR-STRESSED HUMAN UMBILLICAL VEIN
TITLE OF INVENTION: BNDOTHELIAL CELLS
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
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                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/208,572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: CHOATE, HALL, & STEWART
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 424
ATTORIET/AGENT INFORMATION:
NAME: KAPLAN ESQ., WAITEN A.
REGISTRATION NUMBER: 34,199
REFERENCE/CONCET NUMBER: 092662-002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-566
INPORMATION FOR SED ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/237,491
FILING DATE: Concurrently herewith
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MAULEN TYPE: Floppy LLD.
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PC-DOS/MS-DOS
41.0.
TITLE OF INVENTION: ECEF
TITLE OF INVENTION: DERIVATIVES
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SSEE: Pennie & Edmonds
F: 2730 Sand Hill Road
Menlo Park
: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           H: 1 base pairs
nucleic acid
DEDNESS: single
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PRIOR APPLICATION DATA:
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                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                   Boston
                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U. ZIP: 94025
                                                                                                                                                                                                 02109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
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Gaps
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APPLICATION NUMBER: US 08/137,951
FILING DATE: 14-0CT-1993
PLOR APPLICATION DATA:
APPLICATION NUMBER: US 08/197,801
FILING DATE: 17-FEB-1994
ATOMNEY/AGENT INFORMATION:
NAME: Halluin, Albert P.
REGISTRATION NUMBER: 25,227
                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 8135-025
TELECOMMUNICATION IFFORMATION:
TELEPHONE: (415) 854-3660
TELEFAX: (415) 854-3660
TELEFAX: (614) PENNIE
INFORMATION POR SEQ ID NO: 6868:
ENGURNCE CHARACTERISTICS:
LENGTH: 1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Search completed: August 25, 2002, 00:32:24
                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: CDN
IMMEDIATE SOURCE:
CLONE: Z-068903
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(without alignments)
69.294 Million cell updates/sec
                                                                                                                                                              August 24, 2002, 21:19:33; Search time 787.61 Seconds
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1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

4: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*
GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1737567 seqs, 1364410299 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maximum Match 100%
Listing first 45 summaries
                                                                                                           OM nucleic - nucleic search, using sw model
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                US-10-077-383-9
20
                                                                                                                                                                                                                                                                                                                                                                                     IDENTITY_NUC
                                                                                                                                                                                                                                                                      Title:
Perfect score:
Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Searched:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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-	51773,	51773,	51841,	51841,	52001,	52001,	52051,	52051,	52156,	52156,	52157,	52157,	52177,	52177,	52215,	52215,	52240,	52240,	52272,	52272,	52624,	52624,	52642,	52642,	5266R
Description	Sequence						Sequence			Seguence		Seguence								Sequence	Sequence				
91	US-10-027-632-51773	-10-027-632-	US-10-027-632-51841	US-10-027-632-51841	US-10-027-632-52001	US-10-027-632-52001	US-10-027-632-52051	US-10-027-632-52051	US-10-027-632-52156	US-10-027-632-52156	US-10-027-632-52157	US-10-027-632-52157	US-10-027-632-52177	US-10-027-632-52177	US-10-027-632-52215	US-10-027-632-52215	US-10-027-632-52240	US-10-027-632-52240	US-10-027-632-52272	US-10-027-632-52272	US-10-027-632-52624	US-10-027-632-52624	JS-10-027-632-52642	US-10-027-632-52642	10-10-023-632-6360
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Score	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	_
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		10.7. 1 10.027-632-5173 equence 51773, Application US/1002 EMERAL INFORMATION: EMERAL INFORMATION: EMERAL INFORMATION: TITLE OF INVENTION: Identification TITLE OF INVENTION: Identification FILE REFERENCE: 10827-129 CURRENT APPLICATION NUMBER: US/10/ CURRENT APPLICATION NUMBER: US 60/19 PRIOR PILING DATE: 2000-07-12 PRIOR FILING DATE: 2000-07-12 PRIOR FILING DATE: 2000-07-12 PRIOR PAPLICATION NUMBER: US 60/19 PRIOR FILING DATE: 2000-03-29 PRIOR APPLICATION NUMBER: US 60/19 PRIOR PELING DATE: 2000-03-29 PRIOR PELING DATE: 2000-02-24 PRIOR PELING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23 PRIOR PELING DATE: 1999-11-23 PRIOR PELING DATE: 1999-11-23 PRIOR PELING DATE: 1999-11-23 PRIOR PELING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23 PRIOR PELING DATE: 1999-11-23 PRIOR FILING DATE: 1999-11-23	0.0% Similarity 0.0%; O; Conservative		SULT 2 10-027-632-51773/c Sequence 51773, Application US/10027632 GENERAL INFORMATION: APPLICANT: Wang, David G. TITLE OF INVENTION: Identification and
000000000000000000000000000000000000000		10-027-532-5173 6quence 5173, Appli equence 5174, Appli equence 5172, Bernon equence 5100 equence 1008 eq	milarit		(2-51773/c 1773, Appl FORMATION: ": Wang, Da INVENTION:
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APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Deligner in the Human Genome
TITLE OF INVENTION: Polymorphisms in the Human Genome
TITLE OF INVENTION: Polymorphisms in the Human Genome
TITLE OF INVENTION: Polymorphisms in the Human Genome
CURRENT ELINE DATE: 2002-04-30
CURRENT FILINE DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-20
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/195,18
PRIOR APPLICATION NUMBER: US 60/195,18
PRIOR PELLING DATE: 2000-02-24
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APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
CURRENT FILING DATE: 2002-04-30

PRIOR PELLING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR PILLING DATE: 2000-07-29

PRIOR FILING DATE: 2000-02-29

PRIOR FILING DATE: 2000-02-24

PRIOR FILING DATE: 2000-02-24

PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,289
PRIOR APPLICATION NUMBER: US 60/185,389
PRIOR PILLING DATE: 1999-11-23
PRIOR PILLING DATE: 1999-11-23
PRIOR PILLING DATE: 1999-11-23
PRIOR PILLING DATE: US 60/185,389
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 51841
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR APLICATION NUMBER: US 60/146,002
PRIOR PILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/167,363
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PRIOR FILING DATE: 1999-08-09
                                                                                                                                                                                                                                ; Sequence 51841, Application US/10027632 ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 52001, Application US/10027632 GENERAL INFORMATION:
                                                                                                                                                                                                        US-10-027-632-51841/c
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
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TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
                                                                                                                                                   PRIOR AFFLICATION WORDER: US 00/210,000
PRIOR PLING DATE: 2000-07-12
PRIOR PLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-0-20
PRIOR PELICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 1000-03-29
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-09-08
SOFTWARE: PRASESO fOR WINGOWS VERSION 4.0
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR PELLORION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/195,218
PRIOR PELLONG DATE: 2000-03-24
PRIOR PELLONG DATE: 2000-03-24
PRIOR PELLING DATE: 1999-11-23
PRIOR PELLING DATE: 1999-11-23
PRIOR PELLING DATE: 1999-11-23
PRIOR PELLING DATE: 1999-10-28
PRIOR PELLING DATE: 1999-09-28
PRIOR PELLING DATE: 1999-09-28
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                              CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
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US-10-027-632-51841
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US-10-027-632-52051/c
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.13
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
FRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
FRIOR FILING DATE: 2000-03-24
FRIOR FILING DATE: 2000-03-29
FRIOR FILING DATE: 2000-02-24
FRIOR FILING DATE: 2000-02-24
FRIOR FILING DATE: 1200-02-24
FRIOR FILING DATE: 1999-11-23
FRIOR APPLICATION NUMBER: US 60/15/363
FRIOR APPLICATION NUMBER: US 60/16/363
FRIOR PILING DATE: 1999-10-28
FRIOR APPLICATION NUMBER: US 60/16/002
FRIOR APPLICATION NUMBER: US 60/16/002
FRIOR PILING DATE: 1999-09-28
FRIOR PILING DATE: 1999-09-09
FRIOR FILING DATE: 1999-09-09
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Matches 0; Conservative 0; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-04-30
PRIOR PELICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR PELICNG NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
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US-10-027-632-52001/c
                                               ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-52001
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US-10-027-632-52157
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.123
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-29
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1200-02-24
PRIOR FILING DATE: 1200-02-24
PRIOR FILING DATE: 1209-11-23
PRIOR PAPLICATION NUMBER: US 60/157,363
PRIOR FILING DATE: 1999-103-28
PRIOR PAPLICATION NUMBER: US 60/157,363
PRIOR PAPLICATION NUMBER: US 60/156,368
PRIOR PAPLICATION NUMBER: US 60/156,369
PRIOR PAPLICATION NUMBER: US 60/156,369
PRIOR PAPLICATION NUMBER: US 60/156,369
PRIOR PAPLICATION NUMBER: US 60/156,002
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APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION WUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-03-29
PRIOR PELICATION NUMBER: US 60/193, 483
PRIOR PRILING DATE: 2000-03-29
PRIOR PELICATION NUMBER: US 60/16/36
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 1999-09-28
PRIOR PELICATION NUMBER: US 60/16/358
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-09-28
PRIOR FILING DATE: 1999-09-09-28
PRIOR FILING DATE: 1999-09-09-28
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SOFTWARE: FastSEQ for Windows Version 4.0
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US-10-027-632-52156/c
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; ORGANISM: Human
US-10-027-632-52156
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; ORGANISM: Human
US-10-027-632-52156
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0.0%; Score 0; DB 7; Length 1; 0.0%; Pred. No. 0;

Best Local Similarity

Query Match

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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE. 108827.132
CURRENT PAPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-29
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR PAPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 1909-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR RAPLICATION NUMBER: US 60/156,358
PRIOR RAPLICATION NUMBER: US 60/146,002
0; Gaps
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DAME: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
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Matches 0; Conservative 0; Mismatches
0; Mismatches
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PRIOR FILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-03-29
PRIOR PILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185, 218
PRIOR APPLICATION NUMBER: US 60/167, 363
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/167, 363
PRIOR APPLICATION NUMBER: US 60/156, 358
PRIOR FILING DATE: 1999-09-28
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SOFTWARE: FastSEQ for Windows Version 4.0
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Search completed: August 25, 2002, 00:45:45
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TITLE OF INVENTION: Identification and Napping of Single Nucleotide
TITLE OF INVENTION: DOLYMOTPHISMS in the Human Genome
FILE REPERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,6:2
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/218,006
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                                                                                                                                           Query Match 0.0%; Score 0; DB 7; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 0; DB 7; Length 1;
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NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                         ; Sequence 52177, Application US/10027632 ; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-027-632-52177/c
                                                                                ; ORGANISM; Human
US-10-027-632-52157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Human
                                                                                                                                                                                                                                                                                                         US-10-027-632-52177
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                                                                  TYPE: DNA
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APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: David G.

TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REPERENCE: 108827.123
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-03-20
PRIOR FILING DATE: 2000-03-20
PRIOR FILING DATE: 2000-03-20
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1200-11-23
PRIOR FILING DATE: 1999-11-123
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-103-28
PRIOR FILING DATE: 1999-09-28
PRIOR PAPLICATION NUMBER: US 60/166,002
PRIOR PAPLICATION NUMBER: US 60/166,002
PRIOR PILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-08-09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.0%; Score 0; DB 7; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
PRIOR FILING DATE: 2000-04-20
PRIOR PAPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 52215, Application US/10027632; GENERAL INFORMATION:
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US-10-027-632-52215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Human US-10-027-632-52177
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ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-027-632-52215
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SEQ ID NO 52215
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Job time: 12372 sec

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(without alignments)
50.110 Million cell updates/sec
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/ Cgyn_2_6/ptodata/2/pna/US098_COMB.seq:*
/ Cgyn_2_6/ptodata/2/pna/US098_COMB.seq:*
/ Cgyn_2_6/ptodata/2/pna/US099_COMB.seq:*
/ Cgyn_2_6/ptodata/2/pna/US091_COMB.seq:*
/ Cgyn_2_6/ptodata/2/pna/US094_COMB.seq:*
/ Cgyn_2_6/ptodata/2/pna/US094_COMB.seq:*
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/ Cgyn_2_6/ptodata/2/pna/US0998_COMB.seq:*
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/cgn2_6/ptodata/2/pna/US087_COMB.seq:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21979536 seqs, 10817449327 residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Listing first 45 summaries
                                                                                                                                                           OM nucleic - nucleic search, using sw model
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Maximum DB seq length: 200000000
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Maximum Match 10
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Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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2	No.	Score	Match	Match Length	DB	ID	Description
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O	٦	50	100.0	21	7	US-08-309-335A-807	Sequence 807, App
O	7	20	100.0	21	6	US-08-531-241A-807	807,
ບ	m	20	100.0	21	16	US-09-263-959-807	807,
O	4	20	100.0	22	56	US-09-669-187A-908	908
O	ហ	20	100.0	22	37	US-10-017-995-908	908
	9	20	100.0	24	7	US-08-320-507-8	B, App
O	7	20	100.0	24	~	US-08-320-507-9	6
O	æ	20	100.0	24	7	US-08-320-507-10	10,
	σ	20	100.0	24	7	US-08-320-507-11	11,
O	10	20	100.0	24	18	US-09-487-130-1	1,
	11	20	100.0	24	18	US-09-487-130-2	7
O	12	20	100.0	24	18	US-09-487-130-3	'n
O	13	20	100.0	24	18	US-09-487-130-4	4,
	14	20	100.0	24	18	US-09-487-130-5	ς,
O	15	20	100.0	24	18	US-09-487-130-6	9
υ	16	20	100.0	25	35	US-09-956-584-570432	570
O	17	20	100.0	25	62	US-60-234-017-532893	
	18	20	100.0	56	7	US-08-309-335A-538	
	19	20	100.0	26	σ	US-08-531-241A-538	Sequence 538, App
	20	20	100.0	26	16	US-09-263-959-538	
	21	20	100.0	27	23	US-09-605-699-2355	Sequence 2355, Ap
O	22	20	100.0	27	53	US-09-735-363A-6	Sequence 6, Appli
	23	20	100.0	28	~	PCT-US98-07592-1	¥
	24	20	100.0	28	7	US-08-369-881-3317	3317,
	25	20	100.0	28	æ	US-08-408-872-1194	Sequence 1194, Ap
	56	20	100.0	28	12	US-08-842-608A-1	1, A
	27	20	100.0	28	12	US-08-862-610-1	1,
υ	28	20	100.0	28	12	US-08-893-971-2	7
O	29	20	100.0	28	16	US-09-281-481-2	~
ပ	30	20	100.0	28	16	US-09-281-481A-2	~
O	31	20	100.0	28	32	US-09-857-723-181	181

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/cgn2_6/ptodata/2/pna/US6001_COMB.seq:*
/cgn2_6/ptodata/2/pna/US6002_COMB.seq:*
/cgn2_6/ptodata/2/pna/US6003_COMB.seq:*
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us-10-077-383-29.rnpm

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Sequence 807, April cation US/09263959
GENERAL INFORMATION:
APPLICANT: Hood, Leroy E.
APPLICANT: Rowen, Lee
APPLICANT: Koop, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH U
                                                      DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 20; DB 9; Length 21; 100.0%; Pred. No. 7.4e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
                                                                                                         E: Seed and Berry LLP 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/531,241A
FILING DATE: 19-SEP-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY, AGENT INFORMATION:
NAME: MOMASters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 240
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 807:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: McMasters, David D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-MAR-1999
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APPLICANT: Hood, Leroy E. APPLICANT: Koop, Ben F. APPLICANT: Rowen, Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 21 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
                                                  TITLE OF INVENTION: DIA
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and 1
                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                               Washington
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ZIP: 98104-7092
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ZIP: 98104-7092
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CTTY: Seattle
                                                                                                                                                                                                                                                                                                                                             FILING DATE: 19
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                Seattle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-263-959-807/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-531-241A-807
                                                                                                         ADDRESSEE:
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APPLICANT: Koop, Ben F.
APPLICANT: Rowen, Lee
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
CORRESPONDENCE : 1215
CORRESPONDENCE ADDRESS:
                                   Sequence 4, Appli
Sequence 4, Appli
Sequence 46, Appl
Sequence 28, Appl
                                                                                                       Sequence 25724, A Sequence 1242, Ap Sequence 1242, Ap
                   Sequence 4, Appli
                                                                                                                                                             Sequence 364, App
Sequence 60, Appl
Sequence 18866, A
                                                                                                                                                                                                                    Sequence 66, Appl
Sequence 66, Appl
 Sequence 3, Appli
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                                                                                                       8 US-09-704-424-25724

5 US-09-213-6208-11242

6 US-09-213-620C-11242

8 US-09-411-999-364

8 US-09-615-702-18866

PCT-US94-04705-66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5: Seed and Berry
6300 Columbia Center, 701 Fifth Avenue
               US-08-893-971-4
US-09-281-481-4
US-09-281-481A-4
                                                                    US-08-267-803-46
US-08-469-802-28
                                                                                                                                                                                                                                     08-08-055-985-66
 US-07-795-897-3
                                                                                                                                                                                                                                                                                          ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UMBER: US/08/309,335A
19-SEP-1994
                                                                                                                                                                                                                                                                                                                                                            US-08-309-335A-807/c; Sequence 807, Application US/08309335A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 807, Application US/08531241A; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: MOMBSTERS, David D.
REGISTRATION NUMBER: 33,963.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 807:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 agagagagagagagagag 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 21 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
FILING DATE: 19-SEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 2
US-08-531-241A-807/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-309-335A-807
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 STATE:
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us-10-077-383-29.rnpm

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; Sequence 9, Application US/08320507
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Matches 20; Conservative
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
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  SEQ ID NO 908
LENGTH: 22
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                                                   TYPE: DNA
                                                                                             FEATURE:
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TITLE OF INVENTION: Inhibition of Angiogenesis by Nucleic Acids FILE REPRENCE: C1037/7025 (HCL/MAT)

CURRENT APPLICATION NUMBER: US/10/017,935

CURRENT FILING DATE: 2001-12-18

PRIOR PILING DATE: 2000-12-18

PRIOR FILING DATE: 1000-12-14

NUMBER OF SEO ID NOS: 1093

SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                         Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 23); DB 26; Length 22; 100.0%; Pred. No. 7.4e+02; trive 0; Mismarches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                      0; Indels
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| Sequence 908. Application US/09669187A
| Sequence 108. Application US/09669187A
| GENERAL INFORMATION:
| APPLICAMY: Krieg, Arthur M. |
| APPLICAMY: Schetter, Christian | APPLICAMY: Schetter, Christian |
| TITLE OF INVENTION: Immunostimulatory Mucleic Acids | FILE REFERENCE: C1039/7035 (HCL/MAT) |
| CURRENT PELING DATE: 1000-09-25 | PRIOR FILING DATE: 1999-09-25 |
| PRIOR FILING DATE: 1999-09-25 | PRIOR APPLICATION NUMBER: US 60/156,113 |
| PRIOR APPLICATION NUMBER: US 60/156,135 |
| PRIOR APPLICATION NUMBER: US 60/227,436 |
| PRIOR FILING DATE: 2000-08-23 |
| NUMBER OF SEQ ID NOS: 1145 |
| SEQ ID NO 908 | SEC ID NO 908 |
| TENNOW.
                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.0%; Pred. No. 7.4e+02;
Matches 20; Conservative 0; Mismatches 0;
REGISTRATION NUMBER: 33,963
REPERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Synthetic Sequence US-09-669-187A-908
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                                                                                                                 807:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20; Conservative
                                                                                                                                                              LENGTH: 21 base pairs
                                                                                                                                                                               nucleic acid
EDNESS: single
                                                                                                                                                                                                                                   linear
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Best Local Similarity
                                                                                                                                                                                                     STRANDEDNESS:
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0; Gaps
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CORRESPONDENCE ADDRESS: ADDRESSEE: Daniel F. Perez STREET: 703 MCKinney Ave., Suite 316
                                                                                                                              Query Match 100.0%; Score 20; DB 37; Length 22; Best Local Similarity 100.0%; Pred. No. 7.4e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 20; DB 7; Length 24; 100.0%; Pred. No. 7.4e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
OPERATING SYSTEM: Dos 6.0
SOFTWARE: WordPerfect 5.2
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/320,507
                                          OTHER INFORMATION: Synthetic Sequence US-10-017-995-908
                                                                                                                                                                                                                                                                                                                                                                               Sequence 8, Application US/08320507; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 111 TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   33,755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (214) 871-1677
TELEFAX: (214) 817-1083
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 agagagagagagagag 20
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Perez, Daniel F
REGISTRATION NUMBER: 33,7
                                                                                                                                                                                                                                                                       22 AGAGAGAGAGAGAGAGAG 3
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                    nucleic acid
                      APPLICATION NUMBER:
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STREET: /______
TTW: Dallas
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                                       FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-320-507-11
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                                                                                                                                                                                                                                   LENGTH:
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            APPLICANT: Gray
TITLE OF INVENTION: A method for ranking sequences to
TITLE OF INVENTION: select target sequence zones of nucleic acids.
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Daniel F. Perez
STREET: 703 McKinney Ave., Suite 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: A method for ranking sequences to TITLE OF INVENTION: select target sequence zones of nucleic acids. NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 20; DB 7; Length 24; 100.0%; Pred. No. 7.4e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Diskette, 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B: Daniel F. Perez
703 McKinney Ave., Suite 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1110-2012
                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/320,507
FILING DATE: 07-0CT-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/320,507
FILING DATE: 07-0CT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 10, Application US/08320507; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 111
TELECOMDINICATION INFORMATION:
TELEPHONE: (214) 811-1677
TELEFAX: (214) 817-1083
INFORMATION FOR SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Perez, Daniel F
REGISTRATION NUMBER: 33,755
                                                                                                                                                                                                                                                                          OPERATING SYSTEM: Dos 6.0
SOFTWARE: WordPerfect 5.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 agagagagagagagagag 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.09
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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TOPOLOGY: linear
US-08-320-507-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gray
                                                                                                                                                                                      USA
                                                                                                                                                                      Texas
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                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                         75202
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                                                                                                                                                                                      COUNTRY:
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0; Gaps
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TITLE OF INVENTION: A method for ranking sequences to
TITLE OF INVENTION: Select target sequence zones of nucleic acids.
NUMBER OF SEQUENCES: 23
CORRESPONDENCE DEBESS:
                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 20; DB 7; Length 24; 100.0%; Pred. No. 7.4e+02; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Diskette, 3.5 inch, 1.44 Mb storage
IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Daniel F. Perez
STREET: 703 McKinney Ave., Suite 316
                                                                   1110-2012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11110-2012
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/320,507
FILING DATE: 07-0CT-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-320-507-11; Sequence 11, Application US/08320507; GENERAL INFORMATION:
                    NAME: Ferez, Daniel F
REGISTRATION NUMBER: 33,755
REFERENCE/DOCKET NUMBER: 111
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Perez, Daniel F
REGISTRATION NUMBER: 33,755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                              TELEPHONE: (214) 871-1677
TELEFAX: (214) 817-1083
INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 agagagagagagagagag 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: Dos 6.0
SOFTWARE: WordPerfect 5.2
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TELEFAX: (214) 817-1083
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23 AGAGAGAGAGAGAGAG 4
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0
Matches 20; Conservative
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                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                   ; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-320-507-10
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TOPOLOGY: linear
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MEDIUM TYPE: Diskette
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APPLICANT: GRAY, DONALD M.
APPLICANT: HASHEM, GIRAN M.
TITLE OF INVENTION: CONVERSION OF A WATSON-CRICK DNA TO A HOOGSTEEN-PAIRED
TITLE OF INVENTION: DUPLEX.
FILE REFERENCE: 91556/66385CIP
CURRENT APPLICATION NUMBER: 05/99/487,130
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 09/357,424
PRIOR FILING DATE: 1999-07-20
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APPLICANT: HASHEM, GIHAN M.
TITLE OF INVENTION: CONVERSION OF A WATSON-CRICK DNA TO A HOOGSTEEN-PAIRED TITLE OF INVENTION: DUPLEX
FILE REFERENCE: 91556/66385CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: nucleic acid
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CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 09/357,424
PRIOR FILING DATE: 1999-07-20
NUMBER OF SEQ ID NOS: 17
                                                                                                      Sequence 3, Application US/09487130 GENERAL INFORMATION:
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SOFTWARE: PatentIn Ver. 2.1
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Best Local Similarity
                                                                               US-09-487-130-3/c
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US-09-487-130-4
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LENGTH: 24
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Sequence 1, Application US/09487130

Sequence 1, Application US/09487130

GENERAL INFORMATION:
APPLICANT: GRAY, DONALD M.
TITLE OF INVENTION: DUPLEX
TITLE OF INVENTION: DUPLEX
TITLE OF INVENTION: DUPLEX
CURRENT APPLICATION UNBER: US/09/487,130

CURRENT FILING DATE: 2000-01-19

PRIOR PLICATION UNBER: 09/357,424

PRIOR FILING DATE: 1999-07-20
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: GRAY, DOUALD M.
APPLICANT: GRAY, DOUALD M.
TITLE OF INVENTION: CONVERSION OF A WATSON-CRICK DNA TO A HOOGSTEEN-PAIRED TITLE OF INVENTION: DUPLEX
FILE REFERENCE: 91556/6638GIP
CURRENT APPLICATION NUMBER: US/09/487,130
CURRENT FILIG DATE: 1999-07-20
PRIOR FILING DATE: 1999-07-20
PRIOR FILING DATE: 1999-07-20
SOUTHWARE: PALEALITY ON UNBER: 09/357,424
NUMBER: OF SEQ. ID NOS: 17
SEQ. ID NOS: 17
SEQ. ID NOS: 17
SEQ. ID NOS: 17
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Best Local Similarity 100.0%; Pred. No. ...
Matches 20; Conservative 0; Mismatches
       0; Mismatches
                                                   1 agagagagagagagagag 20
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Matches 20; Conservative
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SOFTWARE: PatentIn Ver. 2.1
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Matches 20, Conservative
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APPLICANT: GRAY, DONALD M.
APPLICANT: HASHEM, GIRAN M.
TITLE OF INVENTION: CONVERSION OF A WATSON-CRICK DNA TO A HOGGSTEEN-PAIRED TITLE OF INVENTION: DUPLEX.
FILE REFERENCE: 91556/66385CIP
CURRENT APPLICATION NUMBER: US/09/487,130
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 09/357,424
PRIOR FILING DATE: 1999-07-20
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
              APPLICANT: GRAY, DONALD M.
APPLICANT: GRAY, DONALD M.
APPLICANT: GRAY, DONALD M.
TITLE OF INVENTION: CONVERSION OF A WATSON-CRICK DNA TO A HOOGSTEEN-PAIRED
TITLE OF INVENTION: DUPLEX
FILE REFERENCE: 9156/66385CIP
CURRENT APPLICATION NUMBER: 02/09/487,130
CURRENT PILING DATE: 2000-01-19
PRIOR FILING DATE: 1999-07-20
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTIN VOY: 2.1
SEQ ID NO 5:
LENGTH:: 24
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US-09-487-130-5
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GENERAL INFORMATION:
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LENGTH: 24
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Search completed: August 25, 2002, 00:32:25 Job time: 16383 sec

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US-09-776-479-908/c
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SEQ ID NO 29
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177965,
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69.294 Million cell updates/sec
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12, Appl
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39401, A
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cgn2_6/ptodata/2/pna/USO6_NBW_COMB.seq:*
cgn2_6/ptodata/2/pna/USO6_NBW_COMB.seq:*
cgn2_6/ptodata/2/pna/USO8_NBW_COMB.seq:*
cgn2_6/ptodata/2/pna/USO8_NBW_COMB.seq:*
cgn2_6/ptodata/2/pna/USO9_NBW_COMB.seq:*
cgn2_6/ptodata/2/pna/USIO_NBW_COMB.seq:*
cgn2_6/ptodata/2/pna/USIO_NBW_COMB.seq:*
cgn2_6/ptodata/2/pna/USIO_NBW_COMB.seq:*
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     GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-09-171-209-17
US-09-539-331D-39401
                                                                                                                                                            1737567 seqs, 1364410299 residues
                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-776-479-908
US-10-112-653-877
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Maximum DB seq length: 2000000000
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Match Length DB
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104, App
58757, A
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Haydock, Paul V.
APPLICANT: U'Ren, Jack
APPLICANT: U'Ren, Jack
APPLICANT: Salgene Corporation
ITTLE OF INVENTION: Nucleic Acid Amplification Using an RNA Polymerase and
ITTLE OF INVENTION: DNA/RNA Mixed Polymer Intermediate Products
FILE REPRENEUE: 018048-001710US
CURRENT APPLICATION NUMBER: 105/10/077,383
                                                                                                                                                                                                                                               App
                                                                                                                                                                                                                                                                          Sequence 15613, A
Sequence 21078, A
Sequence 705, App
                                                                                                                                            Sequence 173, App
Sequence 173, App
Sequence 40067, A
Sequence 2057, Ap
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OTHER INFORMATION: a or g at positions 13-20 may be present or absent
                                                                                                                                                                                                                         Sequence 5875'
Sequence 885,
Sequence 1419
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            Sequence 7
Sequence 1
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Sequence 4
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APPLICANT: Petersen, Deanna M.
APPLICANT: Fournon, Yves
IITLE OF INVENTION: Treatment of Asthma and Allergy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 20; DB 6; Length 20; 100.0%; Pred. No. 1.6e+02;
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                       US-09-454-226A-705
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PRIOR FILING DATE: 2001-06-07
NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 908, Application US/09776479; GENERAL INFORMATION:
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Matches 20; Conservative
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; OTHER INFORMATION: Synthetic PCR Primer US-10-078-7578-36
                                                   ORGANISM: Artificial Sequence
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US-10-205-303-265/c
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LENGTH: 50
TYPE: DNA
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APPLICANT: Eerg, Daniel J.
TITLE OF INVENTION: IMMUNOSTIMULATORY NUCLEIC ACID FOR
TITLE OF INVENTION: IMMUNOSTIMULATORY NUN-ALLERGIC INFLAMMATORY DISEASES
FILE REFERENCE: C01039/70060(AWS)
CURRENT APPLICATION NUMBER: US/10/112,653
CURRENT PILING DATE: 2002-03-29
PRIOR APPLICATION NUMBER: US 60/279,642
PRIOR FILING DATE: 2001-03-29
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                                                                                                                                                                                                                                                                                                                                                                                                ch 100.0%; Score 20; DB 5; Length 22; Similarity 100.0%; Pred. No. 1.6e+02; 20; Conservative 0; Mismatches 0; Indels
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APPLICANT: RADER, CAListoph
TITLE OF INVENTION: HUMANIZATION OF MURINE ANTIBODY
FILE REPERRENCE: TSR1 598.0 Con1
CURRENT APPLICATION NUMBER: US/10/078,757B
CURRENT PILING DATE: 2002-02-19
PRIOR FILING DATE: 1997-12-05
PRIOR FILING DATE: 1997-12-05
SOFTWARR F. FALLSCO FOR Mindows Version 4.0
SEQ ID NO 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Synthetic Oligonucleotide US-10-112-653-877
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 877
                                                                                                                         NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 908
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
                                             CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR FILING DATE: 2000-02-03
                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Synthetic Sequence US-09-776-479-908
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Matches 20; Conservative
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.123
CURRENT PELICATION NUMBER: US 60/218,006
PRIOR PELICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR PELICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 1000-02-24
PRIOR PELICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-08-28
PRIOR FILING DATE: 1999-08-28
PRIOR PELING DATE: 1999-08-09
PRIOR REPLICATION NUMBER: US 60/146,002
PRIOR PELING DATE: 1999-08-09
                                                                                    Gaps
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           Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PA120C2
CURRENT APPLICATION NUMBER: US/10/205,303
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Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 3506
Query Match
100.0%; Score 20; DB 7; I
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 20; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0
Matches 20, Conservative
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US-10-205-303-265
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APPLICANT: Brandt, Kevin S.
APPLICANT: Galnes, Petrick J.
APPLICANT: Galnes, Petrick J.
APPLICANT: Ginchcomb, Dan T.
APPLICANT: Wisnewski, Nancy
TITLE OF INVENTION: MCLECULES, PROTEINS AND USES THEREOF
FILE OF INVENTION: MCLECULES, PROTEINS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6947, Application US/09620393B
GENERAL INFORMATION:
APPLICAMY: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTID
TITLE OF INVENTION: THEREBY
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Matches 20; Conservative 0; Mismatches
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CURRENT APPLICATION NUMBER: US/09/620,393B
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 9948
                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/60/319,414
CURRENT FILING DATE: 2002-07-22
NUMBER OF SEQ ID NOS: 2313
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                                                                                                                                          US-60-319-414-2115; Sequence 2115, Application US/60319414; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Ctenocephalides felis
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; OTHER INFORMATION: n = unknown
US-60-319-414-2115
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       1 agagagagagagagag 20
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les 20; Conserva
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: THEREBY
FILLE REPERENÇE: 2750-1068P
CURRENT APPLICATION NUMBER: US/09/620,393B
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 9948
SOFTWARE: Patentin version 3.0
SEQ ID NO 3516
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                                                                                                                   Length 95;
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TILE OF INVENTION: Human Secreted Proteins
FILE REFERENCE: PS550
CURRENT APPLICATION NUMBER: US/10/105,239
CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 15197
Prior Application removed - See File Wrapper or Palm SOFTWARE: Patentin Ver. 2.0
                                                                                                              100.0%; Score 20; DB 7; I
100.0%; Pred. No. 1.5e+02;
ttive 0; Mismarches 0;
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. LOCATION: 1..107
; OTHER: INFORMATION: Ceres Seq. ID 1385553
US-09-620-3938-3516
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; GENERAL INFORMATION:
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; Sequence 8877, Application US/10105299
; GENERAL INFORMATION:
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                                                                                                                                                              20; Conservative
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LOCATION: 1.107
                                                                                                              Query Match
Best Local Similarity
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                      ORGANISM: Human
                                          US-10-027-632-20865
TYPE: DNA
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Sequence 12, Application US/10057940
| GENERAL INFORMATION:
| APPLICANT: Pantoliano, Michael W. | APPLICANT: Salemme, F. Raymond | APPLICANT: Carver, Jr., Theodore, E. | TITLE OF INVENTION: High Throughput Method for Functionally Classifying Proteins | TITLE OF INVENTION: Identified Using a Genomics Approach | FILE REFERENCE: 1503.0310002/JAG/JSO
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APPLICANT: Carver, Jr., Theodore, E.
TITLE OF INVENTION: High Throughput Method for Functionally Classifying Proteins
TITLE OF INVENTION: Identified Using a Genomics Approach
FILE REFERENCE: 1503.0310002/JAG/JSO
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CURRENT FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: 09/190,128
PRIOR FILING DATE: 1998-11-12
PRIOR APPLICATION NUMBER: US 60/065,129
PRIOR FILING DATE: 1997-11-12
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PRIOR APPLICATION NUMBER: US 60/065,129
PRIOR FILING DATE: 1997-11-12
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APPLICANT: Pantoliano, Michael W.
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                                                                                                                  Sequence 7088, Application US/09620393B
GENERAL INFORMATION:
APPLICANT ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: THEREBY
TITLE OF INVENTION: THEREBY
TITLE OF INVENTION: THEREBY
CURRENT APPLICATION NUMBER: US/09/620,393B
CURRENT APPLICATION NUMBER: US/09/620,393B
SOFTWARE: PATENTING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 9948
SOFTWARE: PATENTIN VETSION 3.0
SEQ ID NO 7088
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APPLICANT: Carver, Jr., Theodore, E.
TITLE OF INVENTION: High Throughput Method for Functionally Classifying Proteins
TITLE OF INVENTION: Identified Using a Genomics Approach
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CURRENT APPLICATION NUMBER: US/10/057,940
CURRENT FILING DATE: 2002-05-03
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US-09-620-3938-7088
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PRIOR FILING DATE: 1998-11-12
PRIOR FILING DATE: 1997-11-12
PRIOR FILING DATE: 1997-11-12
NUMBER OF SEO ID NOS: 14
SOFTWARE: PatentIn version 3.0
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    Cgn2_6/ptodata/2/pna/1/5089_COMB. seq:*
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
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/cgn2_6/ptodata/2/pna,/U56000_COMB.seq:* /cgn2_6/ptodata/2/pna,/U56001_COMB.seq:* /cgn2_6/ptodata/2/pna,/U56002_COMB.seq:* /cgn2_6/ptodata/2/pna,/U56004_COMB.seq:* /cgn2_6/ptodata/2/pna,/U56004_COMB.seq:*

Mon Aug 26 10:18:09 2002

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TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MUENCHAU, DARYL, ET AL.
METHOD FOR GENERATING APTAMERS OF
WENTION: NON-PREDETERMINED OR SUBSTANTIALLY
WENTION: NON-PREDETERMINED SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.0%; Score 0; DB 1; Length 1;
                                                     E: Knobbe, Martens, Olson, and Bear
620 Newport Center Dr. Sixteenth Floor
                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/068,298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Marcn
0.0%; Score U; DB 1
0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                              HITACHI.006H
                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US93/00977
FILING DATE: 19930129
                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYRE: Floppy disk
COMPUTER: IBM PC COMPOTEIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICATION NUMBER: 07/659,980
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STREET: LYON & LYON
STREET: 611 West Sixth St.
CITY: Los Angeles
STATE: CA
COUNTRY: CA
                                                                                                                                                                                                                                                                                                                                        NAME: Altman, Daniel E. REGISTRATION NUMBER: 34,115
                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: HISTELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 702: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA to mRNA
                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 714-760-0404
TELEFAX: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: NUCLEIC ACID
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy of
                                        CORRESPONDENCE ADDRESS:
                                                                                         Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION: 435
                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                              USA
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                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HYPOTHETICAL:
                                                                                                              CA
                                                                                                                                             92660
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                                                           ADDRESSEE:
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                                                                                                                              COUNTRY:
                                                                            STREET:
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                                                                      Sequence 895, App
Sequence 982, App
Sequence 1048, App
Sequence 1048, Ap
Sequence 1352, App
Sequence 1352, App
                                                                                                                                                                                              Sequence 51773, A
Sequence 51773, A
Sequence 51841, A
     Sequence 614,
                                        Sequence 836,
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                        Sequence
                                                       Sequence
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US-09-625-102-614
US-09-625-102-836
US-09-625-102-836
US-09-625-102-895
US-09-625-102-895
US-09-625-102-982
US-09-625-102-982
US-09-625-102-1048
US-09-625-102-1352
US-09-625-102-1353
US-09-624-3068-51773
US-09-634-3068-51773
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Knobbe, Martens, Olson, and Bear
STREET: 620 Newport Center Dr. Sixteenth Floor
CITY: Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OMPRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                   US-09-634-306B-51841
                                                                                                                                                                                                                                                                                      ALIGNMENTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 702, Application PC/TUS9300977 ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Altenan, Daniel E.
REGISTRATION NUMBER: 34,115
REFERENCE/DOCKET NUMBER: HITAN
TELEPHONE: 714-760-0404
TELEPHONE: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 702:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO
   4444444444444
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PC
FILING DATE: 19930129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: NUCLEIC ACID
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 7. CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
   00000000000000
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PCT-US93-00977-702
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US-08-068-298-17
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TITLE OF INVENTION: METHOD FOR GENERATING APTAMERS OF
TITLE OF INVENTION: NON-PREDETERMINE) OR SUBSTANTIALLY
TITLE OF INVENTION: NON-PREDETERMINE) SEQUENCES
CORRESPONDENCE 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.0%; Score 0; DB 4; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                       CTHER INFORMATION: where N also may be isoguanine or OTHER INFORMATION: isocytostine US-08-068-298-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PatentIn Release #1.0, Version #1.25
                                                  32,212
BER: 193/051
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APPLICATION NUMBER: US/08/068,298
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REGISTRATION NUMBER: 193/051
TELECOMUNICATION INFORMATION:
TELEPHONE: 213-489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 17, Application US/08068298
; GENERAL INFORMATION:
; APPLICANT: MUENCHAU, DARYL, ET AL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/659,980
FILING DATE: February 21, 1991
ATTORNEY/AGENT INFORMATION:
FILING DATE: February 21, 1991 ATTORNEY/AGENT INFORMATION:
                                NAME: CORSALVI, MARY S.
REGISTRATION NUMBER: 32,212
REFERENCE/DOCKET NUMBER: 193/0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 213-459-1600
TELEFAX: 213-955-0440
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: LYON & LYON
STREET: 611 West Sixth St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
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                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                   LENGTH: 1 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Los Angeles
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                                                                                                                                                                                                                                                                                                                                                                  Sequence 9, Application US/08154792A
GENERAL INFORMATION:
APPLICANT: Kolodier, Richard D.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF INVENTION: and Uses Therefor
NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                Query Match 0.0%; Score 0; DB 4; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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OTHER INFORMATION: where N also may be isoguanine or OTHER INFORMATION: isocytostine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OMPUTER: TBM PC compatible
SOFRAMING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/154,792A
FILING DATE: 17-NOV-1993
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; Sequence 9, Application US/08154792A
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 44824
TELEZOMONIOLATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: 20021 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kolodner, Richard D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34,235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-154-792A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
REGISTRATION NUMBER: 34,2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 1 base pairs
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APPLICATION NUMBER: US/08/154,792A
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Resnick, David S. REGISTRATION NUMBER: 34
                                                                                                                                                                                                                                                                         LENGTH: 1 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                    single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
EDNESS: single
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                        FILING DATE: 17 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF INVENTION: and Uses Therefor
NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF INVENTION: and Uses Therefor
NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                                          ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indel:
                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/154,792A FILING DATE: 17-NOV-1993 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: RESIDENCY, DAVIG S.
REGIETRATION NUMBER: 34,235
REGIETRATION NUMBER: 44824
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24, Application US/08154792A GENERAL INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; MOLECULE TYPE: DNA (genomic) US-08-154-792A-9
                                                                                                                                                                   130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                           Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 1 base pairs
                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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                                                                                                                                                                                                                                   USA
                                                                                                                                                                                       Boston
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Gaps
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APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF INVENTION: and Uses Therefor
NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels Matches 0; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/154,792A
FILING DATE: 17-NOV-1993
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PC-DOS/MS-DOS
                                                  ATTORNEY/AGENT INFORMATION:
NAME: Resolick, David S.
REGISTRATION NUMBER: 34,235
REFERENCE/DOCKET NUMBER: 44824
TELEPHONE: (617) 523-3400
TELEPHONE: (617) 523-6440
TELEFAX: (617) 523-6440
TELEFAX: (617) 523-6440
TELEFAX: COSSI STREE UR
INFORMATION FOR SED ID NO: 24:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 522-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-
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                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA (genomic)
17-NOV-1993
N: 435
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LENGTH: 1 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                            ADDART: 150
STREET: 150
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                                                                                                                                            COUNTRY: US
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                                       1; Indels
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/ Match 0.0%; Score 0; DB 5; Length 1; Local Similarity 0.0%; Pred. No. 0; no Conservative 0; Mismarches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES THEREFOR NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: DAVID G. CONLIN; DIKE, 3RONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: RESDICK, David S.
REGISTRATION NUMBER: 34,234
REFERENCE/DOCKET NUMBER: 44824-CIP-2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/163,449A
FILING DATE: 07-DEC-1993
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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GENERAL INFORMATION:
APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 100
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US-08-163-449A-9/c
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     Query Match
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1; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/163,449A
FILING DATE: 07-DEC-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
REGISTRATION NUMBER: 34,234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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APPLICANT: Reenan, Robert A.G.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: AND USES THEREFOR
TUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES THEREFOR
UNDER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
                                                                                                                                ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-163-449A-24; Sequence 24, Application US/08163449A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                           PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 4482.
TELEPHONE: (617) 523-3400
TELEPHONE: (617) 523-6440
TELEFAX: (617) 523-6440
TELEFAX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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Gaps

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1; Indels

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APPLICANT: Silberstein Ph.D., David S.
APPLICANT: Balcewicz-Sablinska M.D., Maria K.
APPLICANT: Newman Ph.D., Gale W.
TITLE OF INVENTION: CONTROL OF HIV GROWTH WITH ECEF AND
TITLE OF INVENTION: ECEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Silberstein Ph.D., David S.
APPLICANT: Balcawicz-Sablinska M.D., Maria K.
APPLICANT: Newman Ph.D., Gale W.
TITLE OF INVENTION: CONTROL OF HIV GROWTH WITH ECEF AND
                                                                                       Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match 0.0%; Score 0; DB 6; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: CHOATE, HALL, & STEWART
STREET: Exchange Place, 53 State Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              092662-002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/208,572
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; Sequence 10, Application US/08208572
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                       Sequence 10, Application US/08208572 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: CONTROL OF H
TITLE OF INVENTION: ECEF
TITLE OF INVENTION: DERIVATIVES
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kaplan ESq., Warren A.
RECISTRATION NUMBER: 34,199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (617) 227-5020
TELEFAX: (617) 227-566
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: DNA (genomic) US-08-208-572-10
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 1 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                    US-08-163-449A-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: AND USES THEREFOR
TITLE OF INVENTION: AND USES THEREFOR
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
APPLICATION NUMBER: US/08/163,449A
FILING DATE: 07-DEC-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: RESINICA, DAVIG S.
REGISTRATION NUMBER: 34,234
REFERENCE/DOCKET NUMBER: 34,234
REFERENCE/DOCKET NUMBER: 44824-CIP-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-640
                                                                                                                                                        44824-CIP-2
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07-DEC-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
REGISTRATION NUMBER: 34,234
REFERENCE/DOCKET NUMBER: 4482.
TELECOMMUNICATION:
TELEPHONE: (617) 523-3400
                                                                                                                                                                                                                                                                                                                                                                                                                    ; MOLECULE TYPE: DNA (genomic) US-08-163-449A-24
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TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                LENGTH: 1 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
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APPLICANT: Delegeane, Angelo
APPLICANT: Jernigan, Colleen
APPLICANT: Levine, Wendy B.
TITLE OF INVENTION: SHEAR-SPRESSED HUMAN UMBILICAL VEIN
TITLE OF INVENTION: BNDOTHELIAL CELLS
CORRESPONDENCE: 7326
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
0.0%; Score 0; DB 6; Length 1;
Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches 1; Indels
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Ima PC compatible
SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/208,572
TITLE OF INVENTION: ECEF
TITLE OF INVENTION: DERIVATIVES
NUMBER OF SEQUENCES: 14
CORRESPONCE ADDRESS:
ADDRESSEE: CHOATE, 4 STEWART
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: KAPIDAN ESQ., Warren A.
REGISTRATION NUMBER: 34,199
REFERENCE/DOOKET NUMBER: 092662-002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-566
INFORMATION FOR ESQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 1 base pairs
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FILING DATE: Concurrently herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 6868, Application US/08237491; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Pennie & Edmonds
STREET: 2730 Sand Hill Road
CITY: Menlo Park
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear; MOLECULE TYPE: DNA (genomic) US-08-208-572-10
                                                                                                                                                                                                                                  Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                              COMPUTER READABLE FORM:
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                                                                                                                                                                       USA
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                                                                                                                              Boston
                                                                                                                                                         COUNTRY: USF
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                                                                                                                                  REPERENCE/DOCKET NUMBER: 25.227
REGISTRATION NUMBER: 25.227
REPERENCE/DOCKET NUMBER: 8135-025
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 854-3660
TELERAX: (415) 854-3664
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 6868:
SEQUENCE CHARACTERISTICS:
LENGTH: 1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Search completed: August 25, 2002, 00:32:25 Job time: 16383 sec
                                                             APPLICATION NUMBER: US 08/197,801
FILING DATE: 17-FEB-1994
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/137,951
FILING DATE: 14-0CT-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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SEQ ID NO 51773
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Sequence 51773, A
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69.294 Million cell updates/sec
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52157,
52177,
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                                                                                                                                      August 25, 2002, 00:45:46; Search time 787.61 Seconds
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1: /cgn2_6/ptodata/2/pna/?cT_NEW_COMB.seq:*

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7: /cgn2_6/ptodata/2/pna/TSI0_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/TSI0_NEW_COMB.seq:*

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Matches 0; Conservative 0; Mismatches
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FILE OF INVENTION NUMBER: US/10/027,632
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT ELLING DATE: 2002-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR PLILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1000-02-24
PRIOR FILING DATE: 1999-11-33
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-33
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-38
PRIOR FILING DATE: 1999-11-89
PRIOR FILING DATE: 1999-10-80-99
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NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REPERENCE: 108827.128
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT APPLICATION NUMBER: US 60/1430
FRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-29
FRIOR FILING DATE: 2000-02-29
PRIOR FILING DATE: 2000-02-29
PRIOR FILING DATE: 1200-02-24
PRIOR FILING DATE: 1200-02-24
PRIOR FILING DATE: 1209-03-28
PRIOR FILING DATE: 1999-103-28
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
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APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REPERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT PILLIG DATE: 2002-04-30
PRIOR PILLIGATION NUMBER: US 60/218,006
PRIOR PELLING DATE: 2000-07-12
PRIOR PILLING DATE: 2000-04-20
PRIOR FILLING DATE: US 60/198,676
PRIOR FILLING DATE: US 60/193,483
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PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-12-8
PRIOR PLICATION NUMBER: US 60/156,358
PRIOR PALICATION NUMBER: US 60/166,002
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SOFTWARE: FastSEQ for Windows Version 4.0
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SOFTWARE: FastSEQ for Windows Version 4.0
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ORGANISM: Human
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US-10-027-632-52001
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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TITLE OF INVENTION: Polymorphisms in the Human Genome
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PRIOR PRICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 1090-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-09-28
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PRIOR PILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR PLING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PLING DATE: 1999-11-23
PRIOR PRICRATION NUMBER: US 60/156,358
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR APPLICATION NUMBER: US 60/156,002
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR PLING DATE: 1999-09-8
                                                        CURRENT APPLICATION NUMBER: US/10/027,632 CURRENT FILING DATE: 2002-04-30
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                      CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FLING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
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PRIOR APPLICATION UNBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
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ORGANISM: Human
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US-10-027-632-51841
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT PILING DATE: 2002-04-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
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Matches 0; Conservative 0; Mismarches
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CURRENT FILLNE DATE: 2002-04-30
PRIOR FILLNE DATE: 2002-04-30
PRIOR FILLNE DATE: 2000-01-22
PRIOR PILLNE DATE: 2000-04-20
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PRIOR FILLNE DATE: 2000-03-29
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PRIOR FILLNE DATE: 1999-11-33
PRIOR FILLNE DATE: 1999-11-33
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PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
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                                                        ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-52001
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; SEQ ID NO 52001
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TITLE OF INVENTION: identification and Mapping of Single Nucleotide
TITLE OF INVENTION: identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827 129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT PAPLICATION NUMBER: US/01/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-07-20
PRIOR PILING DATE: 2000-03-20
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 1000-01-123
PRIOR FILING DATE: 1999-11-123
PRIOR FILING DATE: 1999-11-123
PRIOR RILING DATE: 1999-11-123
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Matches 0; Conservative 0; Mismatches
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR PELLING DATE: 2000-02-24
PRIOR PELLING DATE: 1999-11-23
PRIOR PELLING DATE: 1999-11-23
PRIOR PELLING DATE: 1999-09-28
PRIOR PELLOATION NUMBER: US 60/156,358
PRIOR PELLOATION NUMBER: US 60/156,358
PRIOR PELLOATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEO ID NOS: 325720
SOFTWARE: FESTSEO for Windows Version 4.0
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52051
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PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
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SERENAL INCOLUTE.
APPLICANT: Warns...
APPLICANT: Warns...
APPLICANT: Warns.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
               1; Indels
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0.0%; Scott v. 0.9;
Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Mismatches
               0; Mismatches
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CURRENT PILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR PELICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-20
PRIOR PLILING DATE: 2000-03-29
PRIOR PELICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR PELICATION NUMBER: US 60/195,218
PRIOR FILING DATE: 2000-02-24
PRIOR PLILOGATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PRING DATE: 1999-10-28
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PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR PALLIANG DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR PLING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/165,318
PRIOR APPLICATION NUMBER: US 60/165,318
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52157
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PRIOR APPLICATION NUMBER: US 60/156,358
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PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
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            0; Conservative
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US-10-027-632-52157
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APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108427.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
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0; Mismatches
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PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR PELING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR PILING DATE: 2000-03-29
PRIOR PILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/165,218
PRIOR APPLICATION NUMBER: US 60/167,363
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SOFTWARE: FastSEQ for Windows Version 4.0
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52156
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PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
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PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
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PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/185,218
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PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
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PRIOR APPLICATION NUMBER: US 60/156,358
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PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
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Best Local Similarity 0.0%,
Matches 0; Conservative
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ORGANISM: Human
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Delymorphisms in the Human Genome
FILE REFERENCE: 108627-132
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
FRIOR FILING DATE: 2000-07-12
FRIOR FILING DATE: 2000-07-12
FRIOR FILING DATE: 2000-07-29
FRIOR FILING DATE: 2000-02-24
FRIOR APPLICATION NUMBER: US 60/185,218
FRIOR APPLICATION NUMBER: US 60/187,363
FRIOR PALING DATE: 1999-11-23
FRIOR FILING DATE: 1999-11-23
FRIOR FILING DATE: 1999-11-23
FRIOR FILING DATE: 1999-11-23
FRIOR FILING DATE: 1999-11-23
FRIOR PALICATION NUMBER: US 60/146,002
FRIOR PAPLICATION NUMBER: US 60/146,002
FRIOR PAPLICATION NUMBER: US 60/146,002
FRIOR FILING DATE: 1999-08-09
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GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Polymorphisms in the Human Genome:
FILE REFERENCE: 108827.129
CURRENT APPLICATION UNIBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
FRIOR FILING DATE: US 60/218,006
PRIOR FILING DATE: US 60/218,006
PRIOR FILING DATE: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/218,006
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NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 52177, Application US/10027632; GENERAL INFORMATION:
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Best Local Similarity 0.0%;
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                                                                                                                                                               ORGANISM: Human
US-10-027-632-52157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-027-632-52177
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                                                                                                                                      TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 52215, Application US/10027632
GENERAL INFORMATION:
FAPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: 2002-04-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.0%; Score 0; DB 7; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2002.02.03.

PRIOR FILING DATE: 2000-07-12

PRIOR FILING DATE: 2000-04-20

PRIOR PELING DATE: 2000-04-20

PRIOR PELING DATE: 2000-04-20

PRIOR PELING DATE: 2000-03-29

PRIOR PAPLICATION NUMBER: US 60/193, 483

PRIOR APPLICATION NUMBER: US 60/185, 218

PRIOR APPLICATION NUMBER: US 60/167, 363

PRIOR FILING DATE: 1999-11-23

PRIOR FILING DATE: 1999-11-23

PRIOR FILING DATE: 1999-11-23

PRIOR FILING DATE: 1999-10-28

PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-11-23
PRIOR PALICATION NUMBER: US 60/167,363
PRIOR PALICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52177
                                                                                                                                                                                                                                          PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 1999-08-09
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Job time: 12373 sec

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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                       OM nucleic - nucleic search, using sw model
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Maximum DB seq length: 2000000000
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18
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58. /cgn2_6/ptodatu2/2/pna/JUS6011_COMB.seq.*
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75. /cgn2_6/ptodatua/2/pna/JUS6033_COMB.seq.*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	ID		-08-309-335A-51	21	US-09-263-959-515	US-09-081-385-40	US-09-589-606-40	US-09-700-354A-40	39-4	US-09-984-198-40	-309-335A	531-241A	US-09-263-959-807	634		-634	¥	US-10-017-995-908	US-08-320-507-8	-320-	-08-320-507	320-50	-08-487-130-	-08-487-130-	-09-487-130	9-487-1	-09-487	-09-487-1	US-09-956-584-462936	-09-956-584-4	-09-956-584-4703	-09-956-584-5704	US-09-956-584-570432
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RESULT 3
US-09-563-959-515
US-09-663-959-515
Sequence 515, Application US/09263959
Sequence 515, Application US/09263959
Sequence 515, Application:
APPLICANT: Hood, Leroy E.
APPLICANT: ROWen, Lee
APPLICANT: KOOP, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UNIVERSE OF SEQUENCES: 1279
                                                    APPLICANT: Rowen, Lee
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH
NUMBER OF SEQUENCES: 1279
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 18; DB 9; Length 18; 100.0%; Pred. No. 1.8e+03;
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                                                                                                                       Seed and Berry LLP
6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Seed and Berry LLP
6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/531,241A
FILING DATE: 19-SEP-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MBER: US/09/263,959
05-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 246
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-6900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 515:
                                                                                                                                                                                                                                                                                                                                                                                                                            33,963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                  NAME: McMasters, David D. REGISTRATION NUMBER: 33,9
                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
              APPLICANT: Hood, Leroy E. APPLICANT: Koop, Ben F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           McMasters, David D.
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
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MEDIUM TYPE: Floppy o
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CLASSIFICATION:
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                                                                                                                                                            Seattle
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STREET: bout
TTY: Seattle
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US-08-531-241A-515
                                                                                                                                                                                              COUNTRY:
                                                                                                                                          STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
                                                                                                                                                                Sequence 98, Appl
Sequence 538, Appl
Sequence 3104, Ap
Sequence 2355, Ap
                                                                                                                                                   Sequence 98, Appl
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Sequence 538, App
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              Sequence 101189
                                              Sequence 502047,
                                                                   507327,
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2 US-60-234-017-532893
0S-60-234-017-532898
US-08-309-335A-338
US-08-531-241A-538
                                     US-60-234-017-502047
                                                                                                                                          US-09-213-620B-98
US-09-213-620C-98
US-09-263-959-538
US-09-605-698-3104
US-09-605-699-2355
US-09-735-363A-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Seed and Berry
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 18; DB 7; I
100.0%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                    ALIGNMENTS
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                                                                                                                                                                                                                                                                                                                                                               Sequence 515, Application US/08309335A
GENERAL INFORMATION:
APPLICANT: Nood, Leroy E.
APPLICANT: Koop, Ben F.
APPLICANT: Rowen, Lee
TITLE ON INVERTION: LARGNOSTIC AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 515, Application US/08531241A
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATORNET JACKILON: 4,3,0 MAME: MCMSTERS, DAVID D. REGISTRATION NUMBER: 33,963
REFERRNET/DOCKET NUMBER: 24005
TELECOMMUNICATION INFORMATION: TELEPHONE: (206) 622-4900
TELEPHONE: (206) 682-6031
INFORMATION FOR SEQ ID NO: 515:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.08;
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Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98104-7092
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US-08-531-241A-515
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE:
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APPLICANT: Gatanaga, T.
APPLICANT: Gatanaga, T.
APPLICANT: Granger, G.A.
TITLE OF INVENTION: Factors Altering Tumor Necrosis
TITLE OF INVENTION: Factor Receptor Releasing Enzyme Activity, and Methods
NUMBER OF SEQUENCES: 154
                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
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            0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US99/10793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/589,606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40, Application US/09700354A; Sequence 40, Application US/09700354A; GENERAL INFORMATION:
                                                                                                                                                                                      Sequence 40, Application US/09589606 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/964,747
FILING DATE: 05-NOV-1997
APPLICATION NUMBER: 60/030,761
FILING DATE: 06-NOV-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 09/081,385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Wu, Frank
REGISTRATION NUMBER: 41,386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                1 agagagagagagagag 18
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                                                                    TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 AGAGAGAGAGAGAG 2
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Matches 18; Conservative
          Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Diskette
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                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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US-09-589-606-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                               US-09-589-606-40/c
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APPLICANT: Granger, G.A.
TITLE OF INVENTION: Factors Altering Tumor Necrosis
TITLE OF INVENTION: Factor Receptor Releasing Enzyme Activity, and Methods
TITLE OF INVENTION: of Use Thereof
NUMBER OF SEQUENCES: 154
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                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 13; DB 16; Length 18; Best Local Similarity 100.0%; Pred. No. 1.8e+03; Matches 18; Conservative 0; Mismarches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: Windows
SOFTWARE: FastsEg for Windows Version 2.0b
SOFTWRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,385
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.425C2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 682-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 515:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: 08/964,747
FILING DATE: 05-NOV-1997
APPLICATION NUMBER: 60/030,761
FILING DATE: 06-NOV-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 40, Application US/09081385
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Wu, Frank
REGISTRATION NUMBER: 41,386
REFERENCE/POCKET NUMBER: 22000
TELECOMMUNICATION:
TELEPHONE: 650-813-5600
TELEFAX: 650-494-0792
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TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette
                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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US-09-081-385-40/C
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                                                                                                                                                                                                                    US-09-263-959-515
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                                                                                                                                                                                                     TOPOLOGY:
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NAME:
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APPLICANT: Granger, G.A.
TITLE OF INVENTION: Factors Altering Tumor Necrosis
TITLE OF INVENTION: Factor Receptor Releasing Enzyme Activity, and Methods
NUMBER OF SEQUENCES: 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                            Factor Receptor Releasing Enzyme Activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
Granger, Gale A. TITLE OF INVENTION: Factors Altering Tumor Necrosis
                                                                                                     ADDRESSEE: BOZICEVIC, FIELD, & FRANCIS, LLP
STREET: 200 MIDDLEFIELD ROAD, #200
CITY: Menlo Park.
                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,09/700,354A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Francis, Carol L.
REGISTRATION NUMBER: 36,513
REFERENCE/DOCKET NUMBER: IRVN-007CIP2
TELECOMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/081,385
FILING DATE: 14-MAY-1998
APPLICATION NUMBER: CT/US99/10793
FILING DATE: 14-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 40:
                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 17-Apr-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                      IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40, Application US/09752639 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 650-327-3400
TELEFAX: 650-327-3231
                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
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INFORMATION FOR SEQ ID NO: 40:
                                                               NUMBER OF SEQUENCES: 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: Windows
                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l agagagagagagagag 18
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Matches 18; Conservative
                                                                                                                                                                                       COUNTRY: USA
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MEDIUM TYPE: Diskette
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APPLICANT: Gatanaga, T.
APPLICANT: Gatanaga, T.
APPLICANT: Granger, G.A.
TITLE OF INVENTION: Factors Altering Tumor Necrosis
TITLE OF INVENTION: Factor Receptor Releasing Enzyme Activity, and Methods NUMBER OF SEQUENCES: 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match 100.0%; Score 18; DB 29; Length 20; Best Local Similarity 100.0%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
SOFTWARE: FASTSEQ for Windows Version 2.0b CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/752,639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 2.0b CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                          22000-20577.21
                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US99/10793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US99/10793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/984,198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 09/081,385
                                                                                                                                                                                                             FILING DATE: 05-NOV-1997
APPLICATION NUMBER: 60/030,761
FILING DATE: 06-NOV-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                          APPLICATION NUMBER: 09/081,385
                                                                                                                                                                                                  08/964,747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 40, Application US/09984198
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: MORRISON & FOERSTER 755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
APPLICATION NUMBER: 08/964,747
FILING DATE: 05-NOV-1997
                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 41,386
                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-813-5600
TELEFAX: 650-494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                       40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Windows
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l agagagagagagagag 18
                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 650-494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19 AGAGAGAGAGAGAG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                  Wu, Frank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 755 PAGE
CITY: Palo Alto
                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94304-1018
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                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                               FILING DATE:
                                                             FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-984-198-40/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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APPLICANT: Hood, Leroy E.
APPLICANT: Koop, Ben F.
APPLICANT: Rowen, Lee
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH U
NUMBER OF SEQUENCES: 1279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hood, Leroy E.
APPLICANT: Rowen, Lee
APPLICANT: Rowen, Lee
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH U
CORRESPONDENCE ADDRESS: 1279
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
PAPLICATION NUMBER: US/08/531,241A
FILING DATE: 19-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                          ADDRESSEE: Seed and Berry LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 18; DB 9; 1
100.0%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                      US-08-531_241A-807/c; Sequence 807, Application US/08531241A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 807, Application US/09263959 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY AGENT INFORMATION:
NAME: MCMSters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 2400:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEPHONE: (206) 682-6931
INPORMATION FOR SEQ ID NO: 807:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seed and Berry LLP
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MEDIUM TYPE: Floppy disk
        1 agagagagagagagag 18
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                              21 AGAGAGAGAGAGAGAG 4
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 21 base pairs
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EDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                        Washington
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US-08-531-241A-807
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GENERAL INFORMATION:
APPLICANT: Hood, Leroy E.
APPLICANT: Koop, Ben F.
APPLICANT: Rowen, Lee
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES: 1215
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 13; DB 36; Length 20; Best Local Similarity 100.0%; Pred. No. 1.8e+03; Matches 18; Conservative 0; Mismarches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 13; DB 7; Length 21; 100.0%; Pred. No. 1.8e+03; Live 0; Mismarches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seed and Berry
6300 Columbia Center, 701 3ifth Avenue
                                                                                                     22000-20577.21
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
APPLICATION NUMBER: 60/030,761
FILING DATE: 06-NOV-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 2400:
TELEPHONE: (206) 622-4900
TELEPAX: (206) 682-691
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 807:
SEQUENCE CHARACTERISTICS:
                                                             NAME: Wu, Frank
REGISTRATION NUMBER: 41,386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: McMasters, David D. REGISTRATION NUMBER: 33,963
                                                                                                  REFERENCE/DOCKET NUMBER: 220
TELECOMUNICOTATION INFORMATION:
TELEPHONE: 650-813-5600
TELEFAX: 650-494-0792
                                                                                                                                                                                                       40:
                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 40
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 agagagagagagagag 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                          linear
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TELEFAX: bov
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US-08-309-335A-807
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US-09-984-198-40
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Gaps

0; Indels

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100.0%; Score 18; DB 24; Length 21; 100.0%; Pred. No. 1.8e+03; ative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICAMT PADISOV, MIKhail, I.
TITLE OF INVENTION: DRUG-CARRIER COMPLEXES AND METHODS OF
TITLE OF INVENTION: USE THEREOF
FILE REFERENCE: 0838.1003-001
CURRENT APPLICATION NUMBER: US/09/634,320
CURRENT FILING DATE: 2000-08-09
PRIOR PILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunostimulatory Nucleic Acids
                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Synthetic Oligonucleotide US-09-634-320-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Synthetic Oligonucleotide US-09-634-320-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Krieg, Arthur M.
APPLICANT: Schetter, Christian
APPLICANT: Vollmer, Jorg
TITLE OF INVENTION: Immunostimulatory Nuclei,
FILE REFERENCE: C1039/7035 (HCL/Mar)
CURRENT PAPLICATION NUMBER: US/09/669,187A
CURRENT FILLING DATE: 2000-09-25
PRIOR APPLICATION NUMBER: US 60/156,113
                                                              PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1999-09-25
PRIOR APPLICATION NUMBER: US 60/156,135
PRIOR FILING DATE: 1999-09-27
PRIOR APPLICATION NUMBER: US 60/227,436
PRIOR FILING DATE: 2000-08-23
               CURRENT FILING DATE: 2000-08-09
PRICR APPLICATION NUMBER: US 60/147,919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-09-25
SER: US 60/156,113
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; Sequence 908, Application US/09669187A
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 6, Application US/09634320 ; GENERAL INFORMATION:
                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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Best Local Similarity
' - h-os 18; Conserva
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                                                                                                                                      SEQ ID NO 10
LENGTH: 21
                                                                                                                                                                                        TYPE: DNA
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100.0%; Score 18; DB 16; Length 21;
Best Local Similarity 100.0%; Pred. No. 1.8e+03;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 18; DB 24; Length 21; 100.0%; Pred. No. 1.8e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/09634320
GENERAL INFORMATION:
APPLICANT: PAPISOV, Mikhail, I.
TITLE OF INVENTION: DRUG-CARRIER COMPLEXES AND METHODS OF TITLE OF INVENTION: USE THEREOF
FILLE REFERENCE: 0838.1003-001
CURRENT APPLICATION NUMBER: US/09/634,320
CURRENT FILING DATE: 2000-08-09
PRIOR APPLICATION NUMBER: US 60/147,919
PRIOR REILIGATION NUMBER: US 60/147,919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10, Application US/09634320; Sequence 10, Application US/09634320; GENERAL INFORMATION: APPLICANT: Papisov, Mikhail, I.
7 APPLICANT: Papisov, Mikhail, I.
7 TILE OF INVENTION: USE THEREOF; TITLE OF INVENTION: USE THEREOF; FILE REFERENCE: 0838.1003-001; CURRENT APPLICATION NUMBER: US/09/634,320
                                              SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Synthetic Oligonucleotide US-09-634-320-5
                                                                                                                                                                                          NAME: MCMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                          APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                        TELEPHONE: (206) 622-4990
TELERAX: (206) 682-6031
INPERMATION FOR SED ID NO: 807:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
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TYPE: nucleic acid
STRANDEDNESS: single
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Matches 18; Conservative
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                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
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US-09-634-320-10/c
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(without alignments)
62.364 Million cell updates/sec
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                                                                                                                                        August 25, 2002, 00:45:46; Search time 787.61 Seconds
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/cgn2_6/ptodata/2/pna/JS06_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/JS06_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/JS08_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/JS09_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/JS10_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/JS10_NEW_COMB.seq:*
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                           OM nucleic - nucleic search, using sw model
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Maximum DB seq length: 2000000000
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Sequence:
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	Description	Seguence 31. Appl	32.	4 . 7	Sequence 29, Appl	90	Sequence 877, App	11, A	29,	11,	25,	7, A	Sequence 36, Appl	Sequence 7, Appli	14	36,	265,		Sequence 8877, Ap		Sequence 2115, Ap	Sequence 6947, Ap	Sequence 7088, Ap	12, Ar	12,	13,	
	ID	US-10-077-383-31	US-10-077-383-32	US-10-011-204-4	US-10-077-383-29	US-09-776-479-908	US-10-112-653-877	US-10-06C-848-11	US-09-997-672-29	US-09-196-447B-11	US-09-724-857-25	US-10-09E-602A-7	US-10-109-624-36	US-10-082-894-7	US-10-16C-362-14	US-10-076-757B-36	US-10-205-303-265	US-10-027-632-20865	US-10-105-299-8877	US-09-62C-393B-3516	US-60-319-414-2115	US-09-62C-393B-6947	US-09-62C-393B-7088	US-10-057-940-12	US-10-057-940-12	US-10-057-940-13	
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Sequence 13, Appl Sequence 75765, A Sequence 178527, Sequence 178533, Sequence 171, App Sequence 171, App Sequence 171, App Sequence 4986, Ap Sequence 16, App Sequence 16, App Sequence 16, App Sequence 16, App Sequence 115, App Sequence 115, App Sequence 115, App Sequence 125, App Sequence 125, App Sequence 125, App Sequence 235, App	Using an RNA Polymerase and ermediate Products Sequence:(XX)-n spacer	Length 18; Indels 0; Gaps 0; ng an RNA Polymerase and diate Products
US-10-057-940-13 US-10-027-632-75765 US-10-027-632-178137 US-10-027-632-178533 US-10-027-632-178533 US-10-027-632-177965 US-10-035-832-323 US-10-035-832-323 US-10-177-632-177965 US-10-177-632-177965 US-10-177-632-177965 US-10-177-240-4986 US-0-171-209-17 US-09-539-3310-40097 US-09-539-3310-40097 US-09-539-3310-185 US-09-963-137-1155 US-09-539-3310-18512 US-09-539-3310-40315	LIGNMENTS 3 nplification Polymer Int 077,383 6,812 Artificial E X = a, Y =	Score 18; DB 6; Le Pred. No. 3.9e+02; Mismatches 0; Mismat
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APPLICANT: Salgene Corporation
APPLICANT: Salgene Corporation
TITLE OF INVENTION: Nucleic Acid Amplification Using an RNA Polymerase and
TITLE OF INVENTION: DNA/RNA Mixed Polymer Intermediate Products
FILE REPRENDENCE: 018048-001710US
CURRENT APPLICATION NUMBER: US/10/077,383
CURRENT PILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: US 60/296,812
PRIOR FILING DATE: 2001-06-07
NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: EXINS, Roger P
TITLE OF INVENTION: Binding assay using binding agents with tail groups
FILE REFERENCE: 0380-P01180US0
                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence:(XY)-n spacer \sim OTHER INFORMATION: Sequence complement, where X = a, Y = g and n = 9 us-10-077-883-32
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CURRENT FILING DATE: 2001-11-08
PRIOR APPLICATION NUMBER: US/08/700,530
PRIOR FILING DATE: 1996-10-23
PRIOR APPLICATION NUMBER: PCT/GB95/00521
PRIOR PILING DATE: 1995-03-10
PRIOR APPLICATION NUMBER: GB 9404709.9
PRIOR FILING DATE: 1994-03-11
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PRIOR FILING DATE: 2001-06-07
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 32
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Matches 18; Conservative
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 4
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                                                                                                                            TYPE: DNA
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; LOCATION: (13)...(20)
; OTHER INFORMATION: a or g at positions 13-20 may be present or absent US-10-077-383-29
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APPLICANT: Berg, Daniel J.
TITLE OF INVENTION: IMMUNOSTIMULATORY NUCLEIC ACID FOR
TITLE OF INVENTION: TREATMENT OF NON-ALLERGIC INFLAMMATORY DISEASES
FILE REFERENCE: C01039/70060(AWS)
CURRENT APPLICATION NUMBER: US/10/112,653
CURRENT FILING DATE: 2002-03-29
PRIOR APPLICATION NUMBER: US 60/279,642
PRIOR FILING DATE: 2001-03-29
                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence:(XY)-n spacer OTHER INFORMATION: Sequence, where X = a and Y = g
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Brazizlar, Robert L.
APPLICANT: Petersen, Deanna M.
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the TITLE OF INVENTION: Treatment of Asthma and Allergy FILE REFERENCE: 01037/7013 (HCL/MAT);
CURRENT APPLICATION NUMBER: US/09/776,479
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PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FASTSEQ for Windows Version 3.0
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SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
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SOFTWARE: PatentIn Ver. 2.1
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Matches 18; Conservative
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                          SEQ ID NO 29
LENGTH: 20
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APPLICANT: Apuya, Nestor R.
APPLICANT: Apuya, Nestor R.
APPLICANT: Tatarinova, Tataian
APPLICANT: Tatarinova, Tataian
APPLICANT: The Regents of the University of California
APPLICANT: Ceres, Inc.
TITLE OF INVENTION: Polynucleotides Use."ul for Modulationg Transcription
FILE REPERENCE: 02307-0118018 195/09/997,672
CURRENT APPLICATION NUMBER: US/09/997,672
PRIOR APPLICATION NUMBER: US/09/997,672
RICH APPLICATION NUMBER: US/09/997,672
RICH APPLICATION NUMBER: US/09/997,672
RICH APPLICATION NUMBER: US/09/997,672
RICH APPLICATION NUMBER: US/09/997,672
                                                                                                                                                                                                                                                                                         0; Gaps
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                                                                                                                                                                                                                                 Query Match 100.0%; Score 13; DB 7; Length 22; Best Local Similarity 100.0%; Pred. No. 3.9e+02; Matches 18; Conservative 0; Mismarches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 13; DB 7; Length 49; 100.0%; Pred. No. 3.9e+02; ttive 0; Mismarches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Univeral primer to poly A tail
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: KLOck, Andrew
APPLICANT: Williams, Deryck Jeremy
APPLICANT: Williams, Deryck Jeremy
APPLICANT: Salmon, Brandy Leigh
APPLICANT: Salmon, Brandy Leigh
TITLE OF INVENTION: NEMATODE MDH-LIKE SEQUENCE
FILE REFERENCE: 12557-002001
CURRENT PAPLICATION NUMBER: US/10/060,848
CURRENT FILING DATE: 2002-06-10
PRIOR APPLICATION NUMBER: US 60/266,037
PRIOR APPLICATION NUMBER: US 60/266,037
PRIOR APPLICATION NUMBER: US 60/266,037
NUMBER OF SEQ ID NOS: 15
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                       ; FEATURE:
; OTHER INFORMATION: Synthetic Oligonucleotide
US-10-112-653-877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-060-848-11; Sequence 11, Application US/10060848; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 29, Application US/09997672; GENERAL INFORMATION:
                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-997-672-29
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; SEQ ID NO 877
; LENGTH: 22
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                                                          TYPE: DNA
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; OTHER INFORMATION: Description of Artificial Sequence:oligonucleotide; OTHER INFORMATION: linker-primer (US-09-997-672-29
                                                                                                                                                                            0; Gaps
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                                                                                                                    Query Match 100.0%; Score 18; DB 5; Length 50; Best Local Similarity 100.0%; Pred. No. 3.9e+02; Matches 18; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNBER: US/09/196,447B
FILING DATE: 19-Nov-1998
CLASSIFICATION: <ur>
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Tripp, Cynthia A.
Frank, Glenn R.
Grieve, Robert B.
TITLE OF INVENTION: NOVEL PARASITIC HELMINTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheridan Ross P.C.
STREET: 1700 Lincoln St., Suite 3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2618-13-3
TELECOMMUNICATION:
TELEPHONE: 303/863-9700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /label- PRIMER
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COCATION: 1.50
CTHER INFORMATION: /label= PRIMED: SEQUENCE DESCRIPTION: SEQ ID NO: 11:
US-09-196-4478-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P22U PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11, Application US/09196447B GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 50 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 303/863-0223
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
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Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
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Query Match
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                                                                                         APPLICANT: Apuya, Nestor R.
APLICANT: Tatainova, Tatiana
APLICANT: Tatainova, Tatiana
APLICANT: Goldberg, Robert B.
APPLICANT: Goldberg, Robert B.
APPLICANT: The Regents of the University of California
TITE. OF INVENTION: Polymucleotides Useful for Modulating Transcription
FILE REFRENCE: 023070-114700US
CURRENT FILING DARE: 2002-05-06
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATENTIN VET. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence:oligonucleotide; OTHER INFORMATION: linker-primer US-09-724-857-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Klock, Andrew P.
APPLICANT: Klock, Andrew P.
APPLICANT: Milliams, Deryck Jeremy
APPLICANT: Sallon, Brandy
TITLE OF INVENTION: NEMATODE GS-LIKE SEQUENCES
FILE REFERENCE: 12557-004001
CURRENT APPLICATION NUMBER: US/10/098,602A
CURRENT FILING DATE: 2002-03-15
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
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Rothberg, Jonathan
Yang, Meijia
US-09-724-857-25; Sequence 25, Application US/09724857; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 7, Application US/10098602A; GENERAL INFORMATION:
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Matches 18; Conservative
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                                                                       APPLICANT: Weterings, Koen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: primer US-10-098-602A-7
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US-10-098-602A-7
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                                                                                                                                                                                                                                                                                                                       SEQ ID NO 25
LENGTH: 50
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Gaps
Knight, James
Kalbfleisch, Theodore
TITLE OF INVENTION: IDENTIFICATION AND COMPARISON OF
PROPEIN-PROTEIN INTERACTIONS THAT OCCUR IN
PROPER PROPER OF THESE
                                                                                                                          POPULATIONS AND IDENTIFICATION OF INHIBITORS OF THESE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 18; DB 7; Length 50; 100.0%; Pred. No. 3.9e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Klock, Andrew P.
APPLICANT: Williams, Deryck Jeremy
APPLICANT: Williams, Deryck Jeremy
APPLICANT: Williams, Deryck Jeremy
APPLICANT: Bradley, John D.
TITLE OF INVENTION: NEMATOBE PGM-LIKE SEQUENCES
FILE REFERENCE: 12557-003001
CURRENT APPLICATION NUMBER: US/10/082,894
CURRENT APPLICATION NUMBER: US 60/271,781
PRIOR FILLING DATE: 2001-02-27
                                                                                                                                                                                                                             ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:

APPLICATION NUBBER: 0S/10/109,624
FILING DATE: 28-MAT-2002
CLASSIFICATION: <a href="https://doi.org/10.10/10/10/10/10/">CLASSIFICATION: <a href="https://doi.org/10/">current</a>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 7934-045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; SEQUENCE DESCRIPTION: SEQ ID NO: 36: US-10-109-624-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION UNBER:
FILING DATE: 13.7UN-1997
APPLICATION WUMBER: 08/663,824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FastSEQ Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 14-JUN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 212-790-9090
TELEFAX: 212-869-8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 7, Application US/10082894; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 50 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                           NUMBER OF SEQUENCES: 122
                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 agagagagagagagag 18
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                                                                                                                                                                                                                                                                              New York
                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                         STATE: NY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
                                                                                                                                                                                                                                                                                                                               COUNTRY:
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GENERAL INFORMATION:
APPLICANT: Williams, Deryck Jeremy
APPLICANT: Williams, Deryck Jeremy
APPLICANT: Salmon, Brandy
APPLICANT: RICek, Andrew
TITLE OF INVENTION: NEWATODE ATP SYNTHASE SUBUNIT E-LIKE SEQUENCES
FILE REFERENCE: 12557-006001
CURRENT APPLICATION NUMBER: US/10/160,362
CURRENT FILING DATE: 2002-05-30
PRIOR PRIOR PILING DATE: 2001-05-31
NUMBER OF SEQ ID NOS: 14
SEQ ID NOS: 14
SEQ ID NO 14
SEQ ID NO 14
SEQ ID NO 14
                                                                                                                                                                                                                                   0; Gaps
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APPLICANT: RADER, Christoph
TITLE OF INVENTION: HUMANIZATION OF MURINE ANTIBODY
FILE REFERENCE: TSRI 598.0 Con1
COUREMY APPLICATION NUMBER: US/10/078,757B
CURREMY TILING DATE: 2002-02-19
PRIOR APPLICATION NUMBER: US 08/986,016
PRIOR APPLICATION NUMBER: US 08/986,016
NUMBER OF SEQ ID NOS: 56
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                 ; OTHER INFORMATION: Univeral primer to poly A tail US-10-082-894-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Synthetic PCR Primer US-10-078-7578-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-078-757B-36; Sequence 36, Application US/10078757B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                             Query Match
Best Local Similarity
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US-10-160-362-14
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SEQ ID NO 36
LENGTH: 50
LENGTH: 50
                                                                     FEATURE:
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Job time: 12373 sec

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45.099 Million cell updates/sec
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/ cgn2_6/ptodata/2/pna/US08_COMB.seq:

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11: /cgn2_6/ptodata/2/pna/USOB3_COMB.seq:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                         OM nucleic - nucleic search, using sw model.
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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18
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44: /ggn2_6/ptddata/2/pna/US6005_COMB.seq:  
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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SUMMARIES	4	ut.	PCT-US93-00977-702	PCT-US93-00977-702	US-08-068-298-17	US-08-068-298-17	US-08-154-792A-9	154-792A-	154-7	-154-792A-	-163-449A-	-163-449A-	163-4	-163-	US-08-208-572-10	-208-2	237-491	-237-4	US-08-296-757B-7984	7578-7		-620-19	-00-306-3	-09-306-3	-09-625-102	-625-102-3	-09-625-102-	-09-625-102-	-09-625-102-	-09-625-102-	-09-65	-09-625-102-	US-09-625-102-614
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/cqn2_6/ptodata/2/pna/US6000_COMB.seq:*
/cqn2_6/ptodata/2/pna/US6001_COMB.seq:*

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TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: MUENCHAU, DARYL, ET AL.
TITLE OF INVENTION: METHOD FOR GENERATING APTAMERS OF
TITLE OF INVENTION: NON-PREDETERMINED OF SUBSTANTIALLY
TITLE OF INVENTION: NON-PREDETERMINED SEQUENCES
                                           SSEE: Knobbe, Martens, Olson, and Bear
: 620 Newport Center Dr. Sixteenth Floor
Newport Beach
                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIALS
OPERATING SYSTEM: PC-DOS/MS-DOS
OFFWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                       HITACHI.006H
                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/00977
FILING DATE: 19930129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/068,298
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17, Application US/08068298 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/659,980
                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E.
REGISTRATION NUMBER: 34,115
                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: HIT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 702: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: LYON & LYON 611 West Sixth St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO
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                   NUMBER OF SEQUENCES: 71 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: NON NUMBER OF SEQUENCES: 26 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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OPERATING SYSTEM:
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                                                                                                                               USA
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PCT-US93-00977-702
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   Sequence 614,
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                                                                                                                                                                                                                                                                                                                                                                                       Sequence 702, Application PC/TUS9300977
GENERAL INFORMATION:
TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
NUMBER OF SEQUENCES: 711
CORRESPONDENCE ADDRESS:
                                                                                                             Sequence
Sequence
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STREET: 620 Newport Center Dr. Sixteenth Floor
CITY: Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.25
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US-09-625-102-895
US-09-625-102-882
US-09-625-102-982
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US-09-625-102-1352
 US-09-625-102-614
US-09-625-102-836
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING COMPATION
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
TELEFAX: 714-760-9502
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E.
REGISTRATION NUMBER: 34,1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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EDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
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HYPOTHETICAL: NO
HOT-US93-00977-702
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17, Application US/08068298
GENERAL INFORMATION:
APPLICANT: MIENCHAU, DARIL, ET AL.
TITLE OF INVENTION: METHOD FOR GENERATING APTAMERS OF
TITLE OF INVENTION: NON-PREDETERMINED OR SUBSTANTIALLY
TITLE OF INVENTION: NON-PREDETERMINED SEQUENCES
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                    0.0%; Score 0; DB 4; Length 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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Best Local Similarity 0.0%; Pred. No. 0;

Matches 0; Conservative 0; Mismatches
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                                                                                  193/051
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/659,980
FILING DATE: February 21, 1991
ATTORNEY/AGENT INFORMATION:
FILING DATE: February 21, 1991 ATTORNEY/AGENT INFORMATION:
                                 NAME: CONSALVI, MARY S
REGIESTRATION NUMBER: 32,212
REFERENCE/DOCKET NUMBER: 193/O
TELECOMMUNICATION INFORMATION:
TELEPHONE: 213-489-160
TELEPHONE: 213-489-160
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
STREET: 611 West Sixth St.
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                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: nucleic acid
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REGISTRATION NUMBER: 33
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                         : 1 base pairs
nucleic acid
                                                                                                                                                                                                                                                 single
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GENERAL INFORMATION:
APPLICANT: Rolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF INVENTION: and Uses Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS &
                                                                                                                                                              1; Indels
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                                                                                                           Query Match 0.0%; Score 0; DB 4; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indel:
OTHER INFORMATION: where N also may be isoguanine or OTHER INFORMATION: isocytostine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.0%; Score 0; DB 5; Length 1;
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OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/154,792A
CIRING DATE: 17-NOV-1993
CLASSIFICATION: 435
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; Sequence 9, Application US/08154792A
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 4482x
TELECHOMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: (617) 523-6440
TELEFAX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kolodner, Richard D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34,235
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; MOLECULE TYPE: DNA (genomic)
US-08-154-792A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: RESNICK, DAVID S.
REGISTRATION NUMBER: 34,2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   130 Water Street
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TYPE: nucleic acid
STRANDEDNESS: single
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       ; OTHER INFO
; OTHER INFO
US-08-068-298-17
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; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-154-792A-24
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GENERAL INFORMATION:
APPLICANT: Rolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: Mismatch Repair Genes, Gene Products,
TITLE OF ENOUNTION: and Uses Therefor
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
                                TITLE OF INVENTION: Mismatch Repair Genes, Gene Products, TITLE OF INVENTION: and Uses Therefor NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                            ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.0%; Score 0; DB 5; Length 1;
                                                                                                                                                                                                                                               COMPUTER: Floppy disk
COMPUTER: LIBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/154 70^**
FILING Name.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 44824
TELECOMMUNICALION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHRRACTERISTICS:
  Reenan, Robert A.G.
Fishel, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34,235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA (genomic) US-08-154-792A-9
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                                                                                                                                                       130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Resnick, David S. REGISTRATION NUMBER: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 1 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                CORRESPONDENCE ADDRESS:
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APPLICANT:
APPLICANT:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: David G. Conlin; DIKE, BRONSTEIN, ROBERTS & ADDRESSEE: CUSHMAN
STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIBLE
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/154,792A FILING DATE: 17-NOV-1993 CLASSIFICATION: 435
APPLICATION NUMBER: US/08/154,792A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 24, Application US/08154792A ; GENERAL INFORMATION:
                     FILING DATE: 17-WOV-1937
CLASSTEICATION: 435
ATTORNEY/AGENT INPORMATION: 435
ATTORNEY/AGENT INPORMATION: 43235
REGISTRATION NUMBER: 34,235
REFERENCE/DOCKET NUMBER: 44824
FELECOMMUNICATION INPORMATION: TELEFAX: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: 200291 STRE UR
INPORMATION FOR SEQ ID NO: 24: SEQUENCE CHARACTERISTICS: LENGTH: 1 base pairs
TELEFAX: 100291 STRE UR
INPORMATION FOR SEQ ID NO: 24: SEQUENCE CHARACTERISTICS: LENGTH: 1 base pairs
TELEFAX: 100291 STRE UR
SEQUENCE CHARACTERISTICS: SEQUENCE CHARACTERISTICS: SINGTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kolodner, Richard D. APPLICANT: Reenan, Robert A.G. APPLICANT: Fishel, Richard
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TELECOMMUNICATION INFORMATION:
TELEFAX: (617) 523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 1 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; MOLECULE TYPE: DNA (genomic) US-08-154-792A-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
REGISTRATION NUMBER: 34,2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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STRANDEDNESS: single
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
REGISTRATION NUMBER: 34,234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; MOLECULE TYPE: DNA (genomic)
US-08-163-449A-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                USA
                                                                                                                         Boston
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US-08-163-449A-24
                                                                                                                                                                COUNTRY:
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                                                                                                                                                STATE:
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                                             0; Gaps
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                                           1; Indels
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Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OFBRATING SYSTEM: PC-FOS_MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATE:
FILING DATE: 07-DEC-1993
                                                                                                                                                                                                                                              APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES, THEREFOR
CORRESPONDENCE ADDRESS: 44
                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: DAVID G. CONLIN; DIKE, ERONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44824-CIP-2
                                                                                                                                                                                                            ; Sequence 9, Application US/08163449A ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9, Application US/08163449A
GENERAL INFORMATION:
APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: RESILCK, DAVIG S.
REGISTRATION NUMBER: 34,234
REGISTRATION NUMBER: 44824
TELECOMMUNICATION INFORMATION:
TELEFHONE: (617) 523-6440
TELEFAX: (617) 523-6440
TELEFX: 200291 STRE UR
INFORMATION FOR SEO ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 1 base pairs
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GENERAL INFORMATION:
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES THEREFOR
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE TITLE OF INVENTION: PRODUCTS, TITLE OF INVENTION: AND USES THEREFOR WOMER OF SEQUENCES: 44 CORRESPONDENCE ADDRESSE: ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: DAVID G. CONLIN: DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/163,449A FILING DATE: 07-DEC-1993 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 44824
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 523-6440
TELEFAX: (617) 523-6440
TELER: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 9:
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APPLICANT: Balcewicz-Sablinska M.D., Maria K.
APPLICANT: Newman Ph.D., Gale W.
ITTLE OF INVENTION: CONTROL OF HIV GROWTH WITH ECEF AND
                                                                                                                                                                                                                                                                                                                                        APPLICANT: Silberstein Ph.D., David S.
APPLICANT: Silberstein Ph.D., David S.
APPLICANT: Balcewicz-Sablinska M.D., Maria K.
APPLICANT: Belcewicz-Sablinska M.D., Maria K.
TITLE OF INVENTION: CONTROL OF HIV GROWTH WITH ECEF AND TITLE OF INVENTION: ECEF
TITLE OF INVENTION: DERIVATIVES
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS: 14
ADDRESSEE: CHOATE, HALL, & STEWART
STREET: Exchange Place, 53 State Street
                                                                                                     1; Indels
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                                                        Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
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APPLICATION NUMBER: US/08/208,572
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 10, Application US/08208572; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                US-08-208-572-10; Sequence 10, Application US/08208572; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: KAPLAN ESTATION NUMBER: 43,199
REFERENCE/DOCKET NUMBER: 0924
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TELEPHONE: (617) 227-5020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , MOLECULE TYPE: DNA (genomic) US-08-208-572-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (617) 227-7566
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
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US-08-163-449A-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kolodner, Richard D.
APPLICANT: Reenan, Robert A.G.
APPLICANT: Fishel, Richard
TITLE OF INVENTION: MISMATCH REPAIR GENES, GENE
TITLE OF INVENTION: PRODUCTS,
TITLE OF INVENTION: AND USES THEREFOR
NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: DAYID G. CONLIN; DIKE, BRONSTEIN, ADDRESSEE: ROBERTS & CUSHMAN STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                44824-CIP-2
                                                                                                                                   44824-CIP-2
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07-DEC-1993
APPLICATION NUMBER: US/08/163,449A FILING DATE: 07-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-163-449A-24/c; Sequence 24, Application US/08163449A; GENERAL INFORMATION:
                                                                                                                            REFERENCE/DOCKET NUMBER: 44824
TELECOMMONICATION INFORMATION:
TELEPRONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: (617) 523-6440
TELEFAX: 20021 STRE UR
INFORMATION FOR SEC ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 44824
TELECHOMUNICATION INFORMATION:
TELEPHONE: (617) 523-3400
TELEFAX: (617) 523-6440
TELEFAX: (617) 523-6440
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34,234
                                                                                                            34,234
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                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Resnick, David S. REGISTRATION NUMBER: 34
                                                                                  NAME: Resnick, David S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 base pairs
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                                                                                                                                                                                                                                                                                                          nucleic acid
EDNESS: single
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                                                                                                            REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                              CLASSIFICATION:
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GENERAL INFORMATION:
APPLICAMT: Sellhamer, Jeffrey
APPLICAMT: Delegeane, Angelo
APPLICAMT: Jernigan, Colleen
APPLICAMT: Levine, Wendy B.
TITLE OF INVENTION: SHEAR-STRESSED HUMAN UMBILICAL VEIN
TITLE OF INVENTION: SHOOTHELIAL CELLS
NUMBER OF SEQUENCES: 7326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.0%; Score 0; DB 6; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE #1.0, Version #1.25
CURRENT APPLICATION DAYA:
APPLICATION NUMBER: US/08/208,572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                    NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
CHARLE, & STEWART
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Kaplan Esq., Warren A.
REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: 092662-002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-5020
TELEPAX: (617) 227-556
INFORMATION FOR SED ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 1 base pairs
TYPE: nucleic acid
STRPE: nucleic acid
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FILING DATE: COGNCURENTLY herewith
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
TITLE OF INVENTION: ECEF
TITLE OF INVENTION: DERIVATIVES
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MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACTIBLE
OPERATING SYSTEM: PC-DOS/MS-
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: 2730 Sand Hill Road
Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                               Floppy disk
                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                                           STREET: EXCHEN
                                                                                                                                                                    USA
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Job time: 16384 sec
                                                                                                                           NAME: Halluin, Albert P.
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 8135-025
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 854-3660
                                                              APPLICATION NUMBER: US 08/197,801 FILING DATE: 17-FEB-1994 ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/137,951
FILING DATE: 14-OCT-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                    TELEPA: (415) 854.3660
TELERA: (415) 843-3694
TELERA: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 6868:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                              LENGTH: 1 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                               IMMEDIATE SOURCE
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MOLECULE TYPE:
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Sequence 51773, A
Sequence 51773, A
Sequence 51841, A
Sequence 52001, A
Sequence 52001, A
Sequence 52001, A
Sequence 52051, A
Sequence 52151, A
Sequence 52157, A
Sequence 52157, A
Sequence 52157, A
Sequence 52177, A
Sequence 52177, A
Sequence 52217, A
Sequence 52217, A
Sequence 52217, A
Sequence 52212, A
Sequence 5264, A
                                                                                                                (without alignments)
62.364 Million cell updates/sec
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                                                                                               August 25, 2002, 00:45:46; Search time 787.61 Seconds
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1: /cgn2_6/ptodata/2/pna/FCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USO6_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

4: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USI0_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USI0_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USI0_NEW_COMB.seq:*
GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
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US-10-027-632-51773
US-10-027-632-51841
US-10-027-632-52001
US-10-027-632-52001
US-10-027-632-52001
US-10-027-632-52051
US-10-027-632-52051
US-10-027-632-52156
US-10-027-632-52157
US-10-027-632-52157
US-10-027-632-52157
US-10-027-632-52177
US-10-027-632-52177
US-10-027-632-522177
US-10-027-632-52644
US-10-027-632-52644
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                                                                                                                                                                                                                                                                                     1737567 seqs, 1364410299 residues
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Maximum Match 100%
Listing first 45 summaries
                                                              OM nucleic - nucleic search, using sw model.
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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	of Single N	Length 1; 1; Indels	f Single Nucleotide
7-632-52668 7-632-52677 7-632-52707 7-632-52714 7-632-52714 7-632-52714 7-632-5272 7-632-52863 7-632-52864 7-632-52864 7-632-52864 7-632-52891 7-632-52891 7-632-52891 7-632-58990 7-632-58990	ents dapping of the Human 32	, DB 7; , 0; atches	Mapping of
US-10-027 US-10-027	LIGN and in 1027, 8,00 8,07 8,07 7,36 5,21 7,36 5,35	Score 0; Pred. No. 0; Misma	C pplication US/10027632 NN: David G.
	1 10	0.0% larity 0.0%; Conservative	3/c Application US IION: 1, David G.
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PRIOR APPLICATION NUMBER: US 60/198, 676
PRIOR FILING DATE: 2000-04-20
PRIOR PELING DATE: 2000-03-29
PRIOR PILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185, 218
PRIOR APPLICATION NUMBER: US 60/167, 363
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-09-28
PRIOR PILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 51841
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                                                                                                                                                                                                                                                                                              Sequence 51841, Application US/10027632 GENERAL INFORMATION:
                                                                                                                                                                                                                                                       US-10-027-632-51841/c
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ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-027-632-52001
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JOURNAL INCRMATION:
JOURNA
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TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108877.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR RILING DATE: 2000-07-12
PRIOR PLICATION NUMBER: US 60/198,676
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR PLICATION NUMBER: US 60/193,483
PRIOR PLICATION NUMBER: US 60/193,483
PRIOR PLICATION NUMBER: US 60/193,383
PRIOR PLILING DATE: 2000-03-29
PRIOR PLILING DATE: 12000-03-29
PRIOR PLILING DATE: 1999-11-23
PRIOR PLILING DATE: 1999-11-23
PRIOR PLILING DATE: 1999-11-23
PRIOR PLILING DATE: 1999-11-23
PRIOR PLILING DATE: 1999-10-28
PRIOR PLILING DATE: 1999-10-28
PRIOR REPLICATION NUMBER: US 60/146,002
PRIOR PLILING DATE: 1999-09-28
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 51841, Application US/10027632; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.0
Best Local Similarity 0.0%
Matches 0; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-51773
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APPLICANT: Wangy David G.

APPLICANT: Wangy David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Dolymorphisms in the Human Genome
FILE REFERENCE: 108827.132
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR RILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/198,528
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR RILING DATE: 1999-10-28
PRIOR PILING DATE: 1999-08-09
PRIOR FILING DATE: 1999-08-09
PRIOR PILING DATE: 1999-08-09
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PRIOR FILING DATE: 2000-03-29
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ORGANISM: Human
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US-10-027-632-52156
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108627.128
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-29
PRIOR FILING DATE: 2000-07-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1200-03-29
PRIOR FILING DATE: 1200-02-24
PRIOR FILING DATE: 1209-03-28
PRIOR FILING DATE: 1999-103-28
PRIOR PAPLICATION NUMBER: US 60/156,363
PRIOR PAPLICATION NUMBER: US 60/156,002
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 10827.129
CURRENT APPLICATION NUMBER: US/10/027,633
CURRENT APPLICATION NUMBER: US/10/0218,006
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-19
PRIOR FILING DATE: 2000-07-19
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Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.)
SEQ ID NO 52001
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                                               ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-52001
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US-10-027-632-52157
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US-10-027-632-52157
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                      TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108827.129 CURRENT APPLICATION NUMBER: US/10/027,632
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
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Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
                                                                                                                        CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR PILING DATE: 2000-04-30
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-03-29
PRIOR PILING DATE: 2000-03-29
PRIOR PILING DATE: 2000-03-24
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-10-28
PRIOR PILICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-08
NUMBER OF SED ID NOS: 335720
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PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-04-30
PRIOR FILING DATE: 2000-04-20
PRIOR PELICATION NUMBER: US 60/198,676
PRIOR PELICATION NUMBER: US 60/193,483
PRIOR PELICATION NUMBER: US 60/193,483
PRIOR PELICATION NUMBER: US 60/185,218
PRIOR PILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/15,363
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
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SOFTWARE: FastSEQ for Windows Version 4.0
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; Sequence 52156, Application US/10027632
; GENERAL INFORMATION:
  APPLICANT: Wang, David G.
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US-10-027-632-52156
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US-10-027-632-52156
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Score 0; DB 7; Length 1;

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Best Local Similarity

Query Match

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Search completed: August 25, 2002, 00:45:46
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TITLE OF INVEWTION: Identification and Mapping of Single Nucleotide
TITLE OF INVEWTION: Debid G.

TITLE OF INVEWTION: Polymorphisms in the Human Genome
TITLE OF INVEWTION: Polymorphisms in the Human Genome
CURRENT APPLICATION NUMBER: US/10/027,6:2
CURRENT FILING DATE: 2000-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR PILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/195,218
PRIOR PILING DATE: 1909-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-10-38
PRIOR PILING DATE: 1999-10-38
PRIOR PILING DATE: 1999-10-38
PRIOR PILING DATE: 1999-10-38
PRIOR PILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SSOFTHARE: FEATSEQ for Windows Version 4.0
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NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52157
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 52177, Application US/10027632 ; GENERAL INFORMATION:
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                                                                                                                                ; ORGANISM: Human
US-10-027-632-52157
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US-10-027-632-52177
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                                                                                                            TYPE: DNA
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APPLICATION DAVIG G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
FILE REFERENCE: 108827.13
FILE REFERENCE: 108827.23
CURRENT APPLICATION NUMBER: US/10/27,632
CURRENT FILING DATE: 2002-04-30
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-07-20
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: US 60/195,218
PRIOR FILING DATE: US 60/195,318
PRIOR FILING DATE: US 60/195,38
PRIOR FILING DATE: 1999-11-23
PRIOR PRILING DATE: 1999-11-23
PRIOR PRILING DATE: 1999-11-23
PRIOR PRILING DATE: 1999-10-28
PRIOR PRILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SSOFTWARE: FEASTEQ for Windows Version 4.0
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PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/193,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-11-23
PRIOR PAPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/166,358
PRIOR APPLICATION NUMBER: US 60/166,002
PRIOR APPLICATION NUMBER: US 60/166,002
PRIOR PELING DATE: 1999-08-28
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52177
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Pizza,M., Scarlato,V., Rappuoll,R., Grandl,G. and Masignani,V.
Neisserial antigens
Patent: WO 9924578-A 25 20-MAY-1899;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Neisserial antigens
Patent: MO 9924678-A 35 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Patent: WO 9924578-A 45 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT); CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
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Sequence 35 from Patent W09924578.
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A96002.1 GI:6779893
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Neisserial antigens
Patent: WO 9924578-A 35 20-MAY-1999;
PIZZA MARRAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
Location/Qualifiers
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Patent: w0 924578-A 25 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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                                                                                                              Neisserial antigens
Neisserial antigens
Patent: WO 9224578-A 45 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
Location/Qualifiers
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Pizza.M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Neisserial antigens
Patent: WO 9224578-A 55 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GIIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
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Sequence 55 from Patent WO9924578.
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Sequence 55 from Patent WO9924578.
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ACCESSION
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Neisserial antigens
Patent: WO 924578-A 55 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza, Mariagna (Tr); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Neisserial antigens.
Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
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Sequence 73 from Patent W09924578.
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Sequence 73 from Patent WO9924578.
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/db_xref="taxon:32644"
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Neisserial antigens
Patent: WO 9924578-A 109 20-MAY-1999;
PIZZA MAKRAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (II); GRANNI GUIDO (IT); MASIGNANI VEGA (IT)
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Patent: WO 9224578-A 109 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANII GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
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Best Local Similarity 100.0%; Pred. No. 0;
Matches 1; Conservative 0; Mismatches
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Sequence 109 from Patent WO9924578.
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Sequence 109 from Patent WO9924578.
A96076
Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches
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/db_xref="taxon:32644"
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Neisserial antigens
Patent: WO 9245/84-A 1 20-MAY-1999;
PIZZA MARIAGRAZIA (IT): SCARLATO VINCENZO (IT): RAPPUOLI RINO (IT);
CHIRON SPA (IT): GRANDI GUIDO (IT): MASIGNANI VEGA (IT)
Location/Qualifiers
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Neisserial antigens
Patent: WO 9924578-A 91 20-MAY-1999;
PIZZA MARRAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Neisserial antigens
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PIZZA MARIAGRAZIA (IT): SCARLAYO VINCENZO (IT): RAPPUOLI RINO (IT);
CHIRON SPA (IT): GRANDI GUIDO (IT): MASIGNANI VEGA (IT)
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result		Query				
No.	Score		Match Length DB ID	DB	OI OI	Description
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m	0	0.0	e	6	AAN80743	Sequence encoding
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5	0	0.0	m	16	AA085491	Plasmid DEZZ318 +h
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7	0	0.0	3	17	AAT33326	CAPL trinucleotide
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KCNQ2; KCNQ3; human; murine; potassium channel; diagnosis; prognosis; behigh familial neonatal epilepsy; BFNB; juvenile myotonic epilepsy; JME; rolandic epilepsy; mutant; treatment; screening; epilepsy; detection; gene therapy; drug screening; primer; ss.
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                                                                                         Human mutant KCNQ3 primer 26.
                     AAX57131 standard; DNA; 1 BP
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                                                                 22-JUL-1999 (first entry)
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                                                                                                               Proteins KCNQ2 and KCNQ3. Detecting mutations in sequences that encode KCNQ2 or KCNQ3, or the loss of one copy of these genes, is used for diagnosis and prognosis of benign familial neonatal epilepsy (BFDE), divenile myotonic epilepsy (AME) or rolandic epilepsy (RE). Cells (or transgent animals) that express wild-type or mutant KCNQ2 or 3 (also the proteins themselves in cell-free form) are used to screen for agents then be used to tream or prevent these forms of epilepsy. Fragments of the encoding nucleic acids are used as probes or primers, either for complete sequences may be used in gene therapy to provide wild-type protein. Antibodies specific for mutant or wild-type proteins are used as diagnostic reagents and for drug screening. The KCNQ2 and 3 proteins are used use useful in rational design of drugs and therapeutically (in
                                                                                                                                                                                                                                                                                                                                                                                                                                          replacement therapies). The forms of epilopsy associated with mutations in KCNQ2 and 3 sequences can now be diagnosed early (before symptoms are manifest), and better treatment options will be available.
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                                                                                                  This invention describes novel human and mouse potassium channel
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Nucleic acid encoding potassium channels KCNQ2 and 3
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Matches 0; Conservative 0; Mismatches
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                                               Claim 65; Page 151; 195pp; English.
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KCNQ2 or KCNQ3, or the loss of one copy of these genes, is used for diagnosis and prognesis of benign familial neonatal epilepsy (BFNE).

Yuvenile myotonic epilepsy (JME) or rolandic epilepsy (RFNE). Cells (or transgenic animals) that express wild-type or mutant KCNQ2 or 3 (also the proteins themselves in cell-free forms) are used to screen for agents that can be used to treat or prevent these forms of epilepsy. Fragments of the encoding nucleic acids are used as probes or primers, either for encoding mutations or for isolation of related sequences, while the protein. Antibodies specific for mutant or wild-type proteins are used complete sequences may be used in gene therapy to provide wild-type proteins are used as diagnostic reagents and for drugs and therapeutically (in replacement therapies). The forms of epilepsy associated with mutations in KCNQ2 and 3 sequences can now be diagnosed early (before symptoms are manifest), and better treatment options will be available.

AAX57074-X57139 are primers used in the method of the invention.
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P-PSDB; AAP80713.
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AAQ85491 standard; cDNA; 3 BP.
                                                                                                                                                                                                                                                                                                                                       Plasmid pEZZ318.thy2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (LASK/) LASKOWSKI M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ANDE/) ANDERSON S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JUL-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9502055-A.
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                                                                                                                                                                                                                                                                          AAQ85491;
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ID AAQ854
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                             A DNA segment with a nucleotide sequence from about 130 to about 918 of the sequence given here is claimed. Also claimed are antibodies which immunoreact with hurfH and the claimed peptides (given in AAP80713). The antibodies may be used in immunoassays for detection of hurfH. The claimed peptides may be used to inhibit the binding of hurfF to coagulation factor VII/VIIa in vivo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A DNA segment with a nucleotide sequence from about 130 to about 918 of the sequence given here is claimed. Also claimed are antibodies which immunoreact with hurFH and the claimed peptides (given in AAP80713). The antibodies may be used in immunoassays for detection of hurFH. The claimed peptides may be used to inhibit the binding of hurF to coagulation factor VII/VIIa in vivo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence encoding complete mature and precursor forms of human tissue factor heavy chain proteins (huTfh & pre-huTfh, respectively).
                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      has gene encoding human tissue factor heavy chain protein and is useful for inhibiting coagulation
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                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                        0.0%; Score 0; DB 9; Length 3;
100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human tissue factor heavy chain (huTFL); immunoassays;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             precursor human tissue factor heavy chain (pre-humeh);
                                                                                                                                                                                             100.0%; Pred. http://weiches
                                                                                                                                        Sequence 3 BP; 0 A; 0 C; 0 G; 0 T; 3 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human tissue factor detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /product-huTFh
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                                                                                                                                                                                                                                                                                                                                                                                                                               10-SEP-1990 (first entry)
Disclosure; ; pp; English.
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                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New DNA segment -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P-PSDB; AAP80713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                          0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     In synthetic analogs of turkey ovomucoid domain protein (6-56) (OWTKY3), given in AAR69818-24, the region immediately adjacent to the reactive site peptide bond is mutated to include the consensus sequence of furin or other serine protease. Polynucleotides encoding such analogs are incorporated into pEZZ318.thy2 and expressed in Escherichia coli RV308 (AFCC 31608).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein inhibitors of serine proteinase(s), e.g. furin, deriv. from turkey ovomucoid third domain - used as laboratory reagents to study the proteinase(s), or as chemotherapeutic agents to treat diseases associated with them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OWTKY3; turkey ovomucoid third domain peptide inhibitor;
furin inhibitor; protease inhibitor; pEZ2318.thy2; probe;
Escherichia coli; ss.
                                                                                                                                                          0; Indels
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                                                                                                Length 3;
                                                                                            DB 9;
Sequence 3 BP; 0 A; 0 C; 0 G; 0 T; 3 other;
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                                                                                  'Match 0.0%; Score 0; DB 9
Local Similarity 100.0%; Pred. No. 0;
les 1; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ85491 standard; cDNA; 3 BP
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AAQ85491;

08-JUL-1994; 09-JUL-1993;

WO9502055-A. 19-JAN-1995.

Synthetic.

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Novel antisense oligonucleotides capable of inhibiting CAPL gene expression may include the trinucleotide GUC (MAT33326, given in 5' to 3' direction) found in codon 14 of CAPL mRNA. These and other antisense oligonucleotides (AAT3337-36) complementary to specific regions of the CAPL gene (see also AAT3335), as well as CAPL-specific ribozymes (AAT33337-40) can be administered to a patient as a means of inhibiting metastatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Engebraaten O, Fodstad O, Hovig E, Maelandsmo GJ;
                                                                                                                                      Agrawal S, Engebraaten O, Fodstad O, Hovig E, Maelandsmo GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic oligo:nucleotide(s) inhibiting CAPL gene expression - useful to inhibit metastatic cancer, partic. osteosarcoma
                                                                                                                                                                                                                                    Synthetic oligo:nucleotide(s) inhibiting CAPL gene expression -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAPL; antisense oligonucleotide; ribozyme; cancer; metastasis;
                                                                                                                                                                                                                                                         useful to inhibit metastatic cancer, partic. osteosarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.0%; Score 0; DB 17; Length 3;
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(NORA-) NORWEGIAN RADIUM HOSPITAL RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.0%; Score 0; DB 17
Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
                                                                                             (NORA-) NORWEGIAN RADIUM HOSPITAL RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3 BP; 0 A; 1 C; 1 G; 1 U; 0 other;
                                                                                                                                                                                                                                                                                                Claim 2; Page 56; 70pp; English.
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                                      95US-0391375.
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  96WO-US02108
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                                                                              (HYBR-) HYBRIDON INC
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16-FEB-1996;
                                      17-FEB-1995;
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                                                                                                                                                         /on Hofe E;
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Von Hofe E;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protein inhibitors of serine proteinase(s), e.g. furin, deriv. from turkey ovomucoid third domain - used as laboratory reagents to study the proteinase(s), or as chemotherapeutic agents to treat diseases associated with them
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAPL; antisense oligonucleotide; ribozyme; cancer; metastasis;
                                                                                                                                      OWNEY3; turkey ovomucoid third domain peptide inhibitor;
furin inhibitor; protease inhibitor; pEZZ318.thy2; probe;
Escherichia coll; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                            93US-0089248.
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                                                         18-AUG-1995 (first entry)
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                                                                                                  Plasmid pEZZ318.thy2.
                                                                                                                                                                                                                                                                                                                                                                                                                                    (LASK/) LASKOWSKI M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-066900/09.
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WO9625499-A1

Synthetic.

AAT33326;

RESULT

Query Match

Matches

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22-AUG-1996.

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Gaps

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1; Indels

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Sequence 3 BP; 0 A; 0 C; 3 G; 0 U; 0 other;

genes of the present invention.

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The present invention relates to human and murine cDNAs from a neurosensory defect associated gene family. The novel cDNAs are mouse tub form I (see AAA94639), mouse tub form II (see AAA94631), human TUB form 6 (see AAA94631), human TUB 1 (see AAA94633), human TUB 1 (see AAA946333), human TUB 1 (see AAA946333), human TUB 1 (see AAA946333), human TUB 1 (see AAA9463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalian proteins expressed in retina and brain, useful for producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibodies and for diagnosing neurosensory defects including cochlear degeneration, peripheral retinal degeneration and cone-rod retinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
Novel antisense oligonucleotides capabble of inhibiting CAPL gene expression may include the trinucleot.ed GUC (AAT33326, given in 5' to 3' direction) found in codon 14 of CAPL mRN. These and other antisense oligonucleotides (AAT3327-36) complementary to specific regions of the CAPL gene (see also AAT33346), as well as CAPL-specific ribozynes (AAT3337-40) can be administered to a patient as a means of inhibiting metasitatic cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3 BP; 0 A; 1 C; 1 G; 1 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.0%; Pred. No.
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Best Local Similarity 0.0%;
--hea 0; Conservative
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01-AUG-1997;
17-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expressing cells and in drug screening assays directed at neurosensory defects. The novel proteins encoded by the present sequence can be used for the treatment of neurosensory degenerative conditions e.g. retinal dystrophies. The present sequence is a probe used to isolate the novel genes of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalian proteins expressed in retina and brain, useful for producing antibodies and for diagnosing neurosensory defects including cochlear degeneration, peripheral retinal degeneration and cone-rod retinal
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0.0%; Score 0; DB 21; Length 3;
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                Pred. No. 0;
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97US-0904699
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to human breast cancer expressed polynucleotides AALD57089) and methods of assessing whether a patient is afflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells.
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tive 0; Mismatches
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24-MAR-2000; 2000US-0193099.
29-MAR-2000; 2000US-0193480.
15-MAY-2000; 2000US-0205230.
09-UUN-2000; 2000US-071171
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The invention relates to human breast cancer expressed polynucleotides (AAL07544-AAL26789) and methods of assessing whether a patient is afflicted with breast cancer by examining the correlation between the arpression of certain markers and the cancerous state of breast cells. The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic
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Human breast cancer expressed polynucleotide 12701.
                                                                   Human; breast cancer; cell marker; cytostatic; ss.
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29-MAR-2000; 2000US-0193480.
15-MAY-2000; 2000US-0205230.
09-JUN-2000; 2000US-0211315.
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                                                                                                                                                                                                                                                      New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
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                                                     94WO-US06884.
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                                                                                                                                             (PHAR-) PHARMAGENICS INC.
                                                                                                                                                                                   Beutel BA, Joesten ME;
                                                                                                                                                                                                                    WPI; 1995-051992/07.
                                                     17-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene therapy.
                                                                                                         07-JAN-1994;
                                                                                         .8-JUN-1993;
                 05-JAN-1995.
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AAQ81664/c
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The sequences given in AAQ81642-95 are oligonucleotides which modulate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the activity of basic fibroblast growth factor (bFGF) by binding to the bFGF protein. Most esp. the sequences given in AAQ81664-67 represent core sequences of which at least one is present in each of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the binding oligos. These oligonucleotides may form a single strand, double strand, a stem-loop structure, a bubble structure, a pseudoknot or a closed, circular structure, bFGF binds to high affinity receptor and low affinity heparin-like molecules on the cell surface. These oligonucleotides bind to bFGF in competition
                                                                                                                                                                                                                     New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.0%; Score 0; DB 16; Length 4; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4 BP; 0 A; 1 C; 2 G; 1 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 25; 44pp; English.
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(PHAR-) PHARMAGENICS INC.
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                                                                     Beutel BA, Joesten ME;
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The sequences given in AAQ81642-95 are oligonucleotides which modulate the activity of basic fibroblast growth factor (bFGF) by binding to the activity of basic fibroblast growth factor (bFGF) by binding to the bFGF protein. Most esp. the sequences given in haAQ81664-67 represent core sequences of which at least one is present in each of the binding oligos. These oligonucleotides may form a single strand, double strand, a stem-loop structure, a bubble structure, a pseudoknot or a closed, circular structure. BFGF binds to high a pseudoknot or a closed, circular structure. BFGF binds to high a finity receptor and low affinity heparin-like molecules on the affinity receptor and heparin. These oligonucleotides may inhibit or chance the activity of bFGF. Particularly, they inhibit concascularisation so they can be used to suppress growth of solid tumours and to reduce the risk of metastasis. They can be used as chipper concepts and the presence of thrombin, or used in
New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
                                                                                                                         Claim 3; Page 25; 44pp; English.
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Query Match 0.0%; Score 0; DB 16; Length 4; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels 1 n 1 1 c 1 q ò

Search completed: August 24, 2002, 22:08:16 Job time: 8849 sec

Sequence 4 BP; 0 A; 2 C; 1 G; 1 U; 0 other;

0; Gaps

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August 24, 2002, 15:57:42; Search time 159.68 Seconds (without alignments) 30.766 Million cell updates/sec
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2: /cgn2_6/ptodata/2/ina/%B_COMB.seq:*
3: /cgn2_6/ptodata/2/ina/%B_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/%B_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/rcTUS_COMB.seq:*
6: /cgn2_6/ptodata/2/ina/rcTuS_COMB.seq:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                      383533 seqs, 122816752 residues
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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20
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Appl Appli		; 0
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2	MEASURING h Floor #1.25	; Length 1; Inc
7.791-213D-62 8.268-679B-7 8.268-679B-7 8.295-743-22 8.602-036A-2 8.602-036A-2 8.293-150A-46 8.293-150A-62 8.293-150A-62 8.293-150A-62 8.293-150A-62 8.642-407A-2 8.642-407A-2 8.642-407A-2 8.642-407A-2 8.642-407A-2 8.873-709-9	MENTS SENT FOR MEA Sixteenth Fl. Version #1. 77	S.
US-08-268 US-08-268 US-08-268 US-08-295 US-08-295 US-08-293 US-08-293 US-08-502 US-08-502 US-08-502 US-08-502 US-08-502 US-08-502 US-08-642 US-08-		Score 0; DB Pred. No. 0; 0; Mismatches
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APPLICATION NUMBER: US/08/268,679B FILING DATE: 30-JUN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07\846,914
FILING DATE: 06-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                         DESCRIPTION: OLIGONUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 086
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 759-4800
TELEFAX: (212) 751-6849
                                                                                                                                                                                     29,323
                                                                                                                                                                                                                                                                         TELEX: 421792
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CRIGINAL SOURCE: N.A.
POSITION IN GENOME: N.A.
US-08-268-679B-8
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NEW YORK
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                                                                                                                                                                     MARIA C.H. LIN
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                                                                                                                                                                                     REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                    UNKNOWN
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                 PCT-US93-00977-702/c
; Sequence 702, Application PC/TUS9300977
; GENERAL INFORMATION:
; TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
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                                                                                                                            ADDRESSEE: Knobbe, Martens, Olson, and Bear
STREET: 620 Newport Center Dr. Sixteenth Floor
CITY: Newport Beach
                                                                                                                                                                                                                                        COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
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APPLICANT: AKHTREUZAMAN; PAUL, ANIKO V.
TITLE OF INVENTION: DE NOVO CELL-FREE
TITLE OF INVENTION: SYNTHESIS PICORNAVIRUS
NUMBER OF SEQUENCES: 8
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345 PARK AVE.
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WORD PERFECT # 5.1
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                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Altenan, Daniel E.
REGISTRATION NUMBER: 34,115
REFERENCE/DOCKET NUMBER: HITA
TELEPHONE: 714-760-0404
TELEPHONE: 714-760-0404
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
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Best Local Similarity 0.0%,
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CLASSIFICATION:
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                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEIC ACID
                                                                                               NUMBER OF SEQUENCES:
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                                                                                                                                                                                                          USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
ZIP: 10154
                                                                                                                                                                                                                          92660
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                                                                                                                                                                                                          COUNTRY:
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0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: WIMMER, ECKARD; WOLLA,
APPLICANT: AKHTERUZZAMAN; PAUL, ANIKO V.
TITLE OF INVENTION: DE NOVO CELL-FREE
TITLE OF INVENTION: SYNTHESIS PICORNAVIRUS
                                                                                                                                      0887-4095 US2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/268,679B FILING DATE: 30-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SYSTEM: PC-DOS/MS-DOS
WORD PERFECT # 5.1
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APPLICATION NUMBER: 07\846,914
FILING DATE: 06-MAR-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 8, Application US/08268679B
Patent No. 5674729
APPLICATION NUMBER: 07\719,761 FILING DATE: 24-JUN-1991 ATTORNEY/AGENT INFORMATION:
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New York
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14603
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                                                                                                                                                                                                                  1 n 1
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                                                                                                                                        Query Match
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APPLICANT: SCOLt, Jeffrey G.
APPLICANT: TOULTA, TARASh1
TITLE OF INVENTION: Cytochrome P4501pr Gene and Its Uses
NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match 0.0%; Score 0; DB 1; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/457,274A
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REGISTRATION NUMBER: 30,727
REFERENCE/DOCKET NUMBER: 19603/240 (D-1519)
TELECOMUNICATION INFORMATION:
TELEPHONE: 716-263-1304
TELEFAX: 716-263-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Nixon, Hargrave, Devans & Doyle
STREET: P.O. Box 1051, Clinton Square
CITY: Rochester
REGISTRATION NUMBER: 29,323
REFERENCE/DOCKET NUMBER: 0887-4095 JS2
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPOTER: TBM PC COMPOTER:
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                DESCRIPTION: OLIGONUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:
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MOLECULE TYPE: DNA (genomic)
                                                                                                TELEX: 421792
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
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CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE: N.A. POSITION IN GENOME: N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 2 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                              SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                      TYPE: NUCLEIC ACID
STRANDEDNESS: SINGL
                                                                                                                                                                                                              UNKNOWN
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MOLECULE TYPE:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Scott, Jeffrey G.
APPLICANT: Tomita, Takashi
TITLE OF INVENTION: Cytochrome P4501pr Gene and Its Uses
NUMBER OF SEQUENCES: 29
                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
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                                                                                                                                                 Query Match 0.0%; Score 0; DB 1; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/457,274A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19603/240 (D-1519)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Nixon, Hargrave, Devans & Doyle
STREET: P.O. Box 1051, Clinton Square
CITY: Rochester
                                                           CHROMOSOME/SEGMENT: Chromosome 1
US-08-457-274A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     POSITION IN GENOME:
CHROMOSOME/SEGMENT: Chromosome 1
                                                                                                                                                                                                                                                                                                                                                 RESULT 6
US-08-457-274A-16/c
: Sequence 16, Application US/08457274A
: Patent No. 5734086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Goldman, Michael L.
REGISTRATION NUMBER: 30,727
REFERENCE/DOCKET NUMBER: 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 716-263-1304
STRAIN: Learn-PyR
DEVELOPMENTAL STAGE: Adult
POSITION IN GENOME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 716-263-1600
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DEVELOPMENTAL STAGE: Adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Musca domestica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 716-203-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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us-10-077-383-9.rni

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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,192
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Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 246102002221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/934,387
FILING DATE: 21-ANG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 28, Application US/08726464B; Patent No. 5932546; GENERAL INFORMATION:
                                                                                                                                                                  3: MORRISON & FOERSTER 755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-813-5600
TELEFAX: 415-494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: GRACEY, NANCY J. REGISTRATION NUMBER: 28,216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BARRETT, Ronald W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_difference
                      LEUNG, LAWRENCE
                                                                                                                                                                                                                                                                                                                  Floppy disk
                                      APPLICANT: VERMAAS, ERIC APPLICANT: TOOLE, JOHN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LATHAM, JOHN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 706141
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
EDNESS: single
                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                       ADDA...
STREET: 755 ...
CITY: PALO ALTO
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
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                                                                                                                                                                                                                                                                         94304
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  APPLICANT:
                      APPLICANT:
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LOCATION: replace[1, "")
OTHER INFORMATION: /note= "This is a biotin-17
OTHER INFORMATION: nucleotide stretch of abasic residues."
                                                                                                                                                                                                                                                                       TITLE OF INVENTION: APTAMERS SPECIFIC FOR BIOMOLECULES AND TITLE OF INVENTION: METHODS OF MAKING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.0%; Score 0; DB 1; Length 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100.0%; Pred. No. 0;
Matches 1; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/934,387
FILING DATE: 21-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/484,192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                             ; Sequence 16, Application US/08484192
; Patent No. 5756291
                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 16, Application US/08484192
; Patent No. 5756291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 24610
TELECOMMUNICATION INFORMATION:
TELEPRONE: 415-4913-5600
TELEPRX: 415-494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 base pairs
                                                                                                                                         APPLICANT: GRIFFIN, LINDA C.
APPLICANT: ALBRECHT, GLENN
APPLICANT: LATHAM, JOHN
APPLICANT: LEUNG, LAWRENCE
APPLICANT: VERMAAS, ERIC
APPLICANT: TOOLE, JOHN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: GRACEY, NANCY J. REGISTRATION NUMBER: 28,216
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EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                         GENERAL INFORMATION:
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                                                           US-08-484-192-16
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APPLICANT: CWIRLA, Steven A.
APPLICANT: CWIRLA, Steven A.
APPLICANT: JOHNSON, Sherril S.
APPLICANT: DUFFIN, David J.
APPLICANT: WAGSTROW, Christopher R.
TITLE OF INVENTION: THROMBOPOIETIN RECEPTOR
TITLE OF INVENTION: APTAMERS SPECIFIC FOR BIOMOLECULES AND TITLE OF INVENTION: METHODS OF MAKING NUMBER OF SEQUENCES: 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
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APPLICANT: Mancharan, Muthiah
APPLICANT: Kanasaki, Andrew
TTPLE OF INVENTION: Anincoxy Modified Oligonucleotides
FILE REFERENCE: 15153824
CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.0%; Score 0; DB 2; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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Patent No. 6127533
GENERAL INFORMATION:
APPLICANT: APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminoxy-Modified Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COCATION: (1)..(2); COCATION: (1)..(2); OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy US-09-016-520-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
                                                   16528A-024100
NAME: Garrett-Wackowski, Eugenia
REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 16528A-024
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/576-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 35, Application US/09016520A
Patent No. 6127533
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                        ; TOPOLOGY: linear
; MOLECULE TYPE: nucleic acid
US-08-726-464B-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                   LENGTH: 2 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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APPLICANT: DOWER, William J.
APPLICANT: UNING, Steven A.
APPLICANT: WISHIN, Steven A.
APPLICANT: WIGHTON, Nicholas C.
APPLICANT: UNFIN, David J.
APPLICANT: WAGSTROW, Christopher R.
TITLE OF INVENTION: THROMBOPOIETIN REJEPTOR
TITLE OF INVENTION: THROMBOPOIETIN REJEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
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ADDRESSEE: TOWNSEND AND TOWNSEND AND CREW
                                                                                                                                                                                                                                                                                                                                    ATTORNEY AGENT INFORMATION:
NAME: Garrett-Wackowski, Eugenia
REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 16528A-024.100
TELECOMMUNICATION INFORMATION:
TELECHONE: 415/576-0200
TELEFAX: 415/576-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
ADDRESSEE: TOWNSEND AND TOWNSEND AND CREW
                        Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Two Embarcadero Center, 8th Floor
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Matches 0; Conservative 0; Mismatches
                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/726,464B
FILING DATE: 04-0CT-1996
CLASSIFICATION: 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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; Patent No. 5932546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: nucleic acid US-08-726-464B-28
                                                                                                                                                             Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 2 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                         San Francisco
California
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                                                                                                                                                                                                       OPERATING SYSTEM:
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APPLICANT: Manoharan, Muthiah
APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
TITLE OF INVENTION: Aninoxy-Modified Oligonucleotides And Methods For
                                                                                                                                                                                                                                                                                                             COTHER INFORMATION: 2'-dimethylaminooxyethyl thymidine (T-2'-DMAOE) COTHER INFORMATION: Description of Artificial Sequence: No. 6172209e1 COTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
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                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
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BER: 09/016,520
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; Patent No. 6194598
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
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NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                          FILE REFERENCE: ISIS2955
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US-09-477-902-35
                                                                                                                                                                                                                            SEQ ID NO 35
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APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Rawasaki, Andraw P
TITLE OF INVENTION: Aminoxy-Modified Oligonucleotides And Methods For
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , LUCATION: (1)..(2); dimethylaminooxyethyl thymidine (T-2'-DMAOE); OTHER INFORMATION: Description of Artificial Sequence: No. 6172209e1; OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                          : LOCATION: (1)..(2)
; CTHEN INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
(MS-09-016-520-35
             CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER PELLICN NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
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; Patent No. 6172209
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                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                 INFORMATION: Sequence
                                                                                                                        SOFTWARE: Patentin Ver. 2.1
    FILE REFERENCE: ISIS2824
                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
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US-09-130-973-35/c
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TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides FILE REFERENCE: ISIS2824 CURRENT APPLICATION NUMBER: US/09/477,902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           h NAME/KEY: misc_feature
   LOCATION: (1)..(2)
   OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
   US-09-477-902-35
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Search completed: August 24, 2002, 21:57:39 Job time: 21597 sec

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56.110 Million cell updates/sec
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMAR1:ES

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1 10 BE401958 1 10 BE402349 1 10 BE402351 1 10 BE402351 1 10 BE402351 1 10 BE402374 1 10 BE402520 1 10 BE402520 1 10 BE402520 1 10 BE402580 1 10 BE402580 1 10 BE402580 1 10 BE402580 1 10 BE402580 1 10 BE402795 1 10 CNSO03FA 1 12 CNSO03FE 1 12 CNSO03FE 1 12 CNSO03FE	ALIGNMENTS RNA; EST; 1 BP.	reated) ast updated, Vers DKFZp564L2071_s1 tag. rdata; Craniata; arrhini; Hominida	L., Brandt P., Mewe) to the EMBL/GenB 18a D-82152 Martin	equenced by GBF w the German Genom at the RZPD in Be Ressourcenzentru ERMANY; Email: cl /Qualifiers "taxon:9606" "taxon:9606" NFTSp564L2071" Ib="564 (synonym:
	17 1 H3M01740 standard; AL037411; AL037411.1	2-MAR-1999 (Rel. 5; 2-MAR-1999 (Rel. 5; 0mo sapiens mRNA; 1 ST; expressed seque ST; expressed seque omo sapiens (human ukaryota; Metazoa; utheria; Primates;		Clone from S. Wiemann, ssequencing consortium of No II sequence available a Plasse contact the RZPD: Berlin-Charlottenburg, G Key Location, Source Above Abo
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HSM003869 standard; RNA; EST; 1 BP.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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Homo sapiens (human)
Bukaryota; Metazoa: Chordata; Craniata; Vertebrata; Buteleostoml; Mammalia;
Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                          Duesterhoeft A., Lauber J., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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0.0%; Pred. No. 0;
ive 0; Mismatches 1; Indels
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12-MAR-1999 (Rel. 59, Created)
12-MAR-1999 (Rel. 59, Last updated, Version 1)
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ri sequence also available
This clone is available at the RZPD in Berlin
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Best Local Similarity 0.0%;
Matches 0; Conservative
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Gaps

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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ID HSM005008 standard; RNA; EST; 1 BP.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                          Duesterhoeft A., Lauber J., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                rl sequence also available
This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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                                                                                                                                                                                                                                                                                                                                                              /tissue_type="testis"
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AL046265;

HSM011115

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Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="44 (synonym: htes3). Vector pSportl; host
DH10B; sites Not! + Sal!"
/dev_stage="adult"
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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                                                                                                                                                                      Koehrer K., Beyer A., Mewes W., Gassenhuber J., Wiemann S.;
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                                                                                                                                                                                                             Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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This clone is available at the RZPD in Berlin
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Matches 0; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/clone="DKF2p434G017"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="testis"
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EST; expressed sequence tag.
                                           Homo sapiens (human)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone="DKFZp434G017"
/clone_lib="434 (synonym: htes3). Vector pSportl; host
DH10B: sites NotI + SalI"
/dev_stage="adult"
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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0.0%; Score 0; DB 2; Length 1;
Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches 1; Indel:
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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DH108: sites Not! + Sall/Mlu!"
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                                                                                                                                                                                                                                                                                                 Homo sapiens mRNA; EST DKFZp586P0817_rl (from clone DKFZp586P0817)
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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      Query Match 0.0%; Score 0; DB 2; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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12-WAR-1999 (Rel. 59, Last updated, Version 1)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone_lib="434 (synonym: htes3). Vector pSport1; host pH10B: aites NotI + SalI"
/dev_stage="adult"
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DH10B; sites NotI + SalI"
/dev_stage="adult"
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email clone@rzpd.de
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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ID HSM011270 standard; RNA; EST; 1 BP.
                                             Location/Qualifiers
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stage subtraction cDNA library (hasp)"

/dev_stage="Seedling roots of scots pine were infected for 6 days with H. annosum"
/note="Vector: pT-Adv; Site_1: EcoRI; The subtractive hybridization cDNA library was constructed from scots pine roots infected for 6-days with mycelia of Heterobasidon annosum (FPS)."

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Anote—"Vector: pT-Adv; Site_1: EcoRI; The subtractive hybridization cDNA library was constructed from scots pine roots infected for G-days with mycelia of Heterobasidon annosum (FP5)."

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/clone="hip="Heterobasidion annosum - Scots pine infection
/clone_lib="Heterobasidion annosum - Scots pine infection
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Dept. of Forest Mycology & Pathology
Swedish University of Agriculture, Box 7026,S-750 07, Uppsala,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Pinus sylvestris/Heterobasidion annosum"
/db_xref="taxon:169015"
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Pinus sylvestris/Heterobasidion annosum
Eukaryota; mixed EST libraries.
( bases 1 to 1)
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Matches 1; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.0%; Score 0;
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Location/Qualifiers
                         Location/Qualifiers
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Seq primer: T7 primer
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Fax: +46 18 30 92 45
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hasp002xj20f Heterobasidion annosum - Scots pine infection stage
subtraction cDNA library (hasp) Pinus sylvestris/Heterobasidion
annosum cDNA clone hasp002xj20f, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              //organism="Homo sapiens"
//organism="Homo sapiens"
//clone="KKFZp586F0817"
//clone_lib="866 (synonym: hutel). Vector pSportl; host
//dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Fred O. Asiegbu
Dept. of Forest Mycology & Pathology
Swedish University of Agriculture, Box 7026,S-750 07, Uppsala,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                     Homo sapiens mRNA; EST DKFZp586P0817_r1 (from clone DKFZp586P0817)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        No sl sequence available
This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                                                                                                    Koehrer K., Beyer A., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Clone from S. Wiemann, sequenced by BMFZ within the cDNA sequencing consortium of the German Genome Project
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Eukaryota; mixed EST libraries.
   12-MAR-1999 (Rel. 59, Created)
12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Fax: +46 18 30 92 45
Email: Fred.Asiegbu@mykopat.slu.se
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BI817896 1 bp mRNA linear EST 04-OCT-2001 G3-023 Axolotl Lambda Zap Library Ambystoma mexicanum cDNA similar to Putative heterochromatin-like protein 1, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fmail: srvoss@lamar.colostate.edu
Single pass sequence from 5' end. Low quality sequence was trimmed
from the ends (PHRED error rate = 5%). Trace file available:
                                                                                                                                                                                                                                                                                                                    Ambystoma mexicanum
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibla; Batrachia; Caudata; Salamandroidea; Ambystomatidae;
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Matches 0; Conservative 0; Mismarches 1; Indels
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/db_xref="taxon:8296"
/clone_lib="Axolot! Lambda Zap Library"
/tissue_type="Regenerating forelimb"
/dev_stage="Medium-bud blattema"
1 c 0 g 0 t
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Location/Qualifiers
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Fort Collins, CO 80523, USA
Tel: 970 491 4869
Fax: 970 491 0649
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BI817896.1 GI:15951508
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Department of Biology
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A51708 Sequence 14
AR167586 Sequence
AR167587 Sequence
             A64735 Sequence 1
A64736 Sequence 2
A64738 Sequence 4
A64739 Sequence 5
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AR167592 Sequence
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AR122341 Sequence
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AR128994 Sequence
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AX115977 Sequence
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Matson. S.S., Coassin, P.J., Rampal, J.B. and Caskey, C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5981185-A 72 09-NOV-1999;
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59;
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                                        August 24, 2002, 21:54:43; Search time 2108.3 Seconds (without alignments) 198.516 Million cell updates/sec
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                                                                                                                               3595312
     GenCore version 4.5
Copyright (c) 1993 - 2007 Compugen Ltd.
                                                                                                                  1797656 seqs, 10463268293 residues
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Listing first 45 summaries
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Sequence 38 Sequence 5

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/organism="unidentified"
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Sequence 4 from Patent W09729116.
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                                                                                                     Reese, C.B. and Rao, M.V.
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               A64736.1 GI:4530772
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1 (bases 1 to 21)
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[ (bases 1 to 20)
Matson, R.S., Coassin, P.J., Rampal, J.B. and Caskey, C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5981185-A 93 09-NOV-1999;
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Reses_C.B. and Rao,M.V.
RUDHUR CONFAINING DINUCLEOTIDE PHOSPHORAMIDITES
Patent: WO 9729116-A 1 14-AUG-1997;
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Sequence 93 from patent US 5981185.
AR084604
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Sequence 1 from Patent WO9729116.
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      UNIVERSITY OF IOWA RESEARCH FOUNDATION (US) ; Coley Pharmaceutical
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Gray, D.M. and Clark, C.L.
Method for selectively ranking sequences for antisense targeting
Patent: US 5856103-A 8 05-JAN-1999;
Location/Qualifiers
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Gray.D.M. and Clark,C.L.
Wethod for selectively ranking sequences for antisense targeting
Patent: US 5856103-A 9 05-JAN-1999;
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Sequence 9 from patent US 5856103.
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Reses.C.B. and Rao,M.V.
DINUCLEUTR CONTAINING DINUCLEOTIDE PHOSPHORAMIDITES
Patent: WO 9729116-A 5 14-AUG-1997;
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1 (bases 1 to 22)

Krieg, A.M., Schetter, C. and Vollmer, J.C.
Immunostimulatory nucleic acids
Patent: Wo 0122972-A 908 05-APR-2001;
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Sequence 908 from Patent WO0122972.
AX104716
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Sequence 5 from Patent W09729116.
A64739
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Unclassize 1 to 24)
Gray,D.M. and Clark,C.L.
Method for selectively ranking sequences for antisense targeting Patent: US 5856103-A 10 05-JAN-1999;
Location/Qualifiers
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Gray, D.M. and Clark, C.L.
Method for selectively ranking sequences for antisense targeting
Patent: US 5856103-A 11_05-JAN-1999;
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1 (bases 1 to 24)
Gray, D.M. and Clark, C.L.
Apparatus and method for selectively ranking sequences for
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Patent: US 6183966-A 10 06-FEB-2001;
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Patent: US 6183966-A 9 06-FEB-2001;
Location/Qualifiers
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Location/Qualifiers
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SNS Gray, D.M. and Clark, C.L.
Apparatus and method for selectively ranking sequences for antisense targeting antisense targeting
RNAL Patent: US 6183966-A 11 06-FEB-2001;
Location/Qualifiers
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AR128996.1 GI:14116658
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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Pred. No. 1s the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

CHMMADIEC

SOMMANIES			Description		14	583 Phosphorothioate o			10		zarger sequence #2			
300			ID		AAT86	AAT86	AA033	AAF99	AA033511	AAXOO	SCHOOL A	2000	AAX00525	AAX00527
			DB	:	18	18	13	22	13	20	2 6	9	50	20
			Match Length DB ID		21	21	22	22	23	24		7	24	24
	æ	Query			100.0	100.0	100.0	100.0	100.0	100.0	1001	2	100.0	100.0
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RESUL AAT86 ID	ESULT 1 AT86582/C D AAT8658	7	standard;	i; DNA;	21	BP.		
X Y ?	AAT8658	182;						
Y L X	25-MAR	8-1998	8 (first	rst entry)	ry)			
DE	Phosp	orot	Phosphorothioate	oligonucleotide	nclec	tide #1.		
X X X X X X X	Phosphoro thioester ss.	oroti	Phosphorothioate thioester; DNA sy SS.	e oligonucleotide; synthesis; antisen	uclec s; an	OΩ	dimeric phosphoramidite e oligonucleotide; gene	ite synthon; ene therapy;
×		,						
FT	Key misc_c	diffe	erence]	cocation 1.21 /*tag=	n/Qua a Phos	Location/Qualifiers 121 /*tag= /*hosphorothioate nucleotides (1 and	linkages betw 12, 3 and 4	een alternate etc.)"
X Z	WO9729116	116-	-A1.					
X G.	14-AUG	-AUG-199	7.					
AY Y	06-FEB-1	3-1997		97WO-GB00327	0327.			
PR S	06-FEB-1	3-1996		96GB-0002326	2326.			
44	,							

Gaps

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was prepared by solid phase synthesis. The method comprises adding at least one dimeric phosphoramidite synthon, optionally having a protected thiosester group in its internucleotide link, during the synthesis cycle. These novel dimeric phosphoramidite synthons are used as antiseanse molecules for inhibition of gene expression. The method gives increased
                     The present sequence represents a phosphorothioate oligonucleotide which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      screening a library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TO)15 oligouncleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites in the bovine genome is estimated at >100. One sequence information for ca 230 such bovine microsatellites is summarised in the specification and indexed herein (see below).
                                                                                                                                                                 yields of the phosphorothiate oligonucleotide (since fewer cycles are needed) and facilitates separation of impurities (greater difference in size compared with use of monomeric synthons).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            were used to generate the required PCR primers for in vitro amplification of the corresp microsatellite (using the program opprentum). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequences upstream and downstream of the microsatellite sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.0%; Score 20; DB 18; Length 21;
Best Local Similarity 100.0%; Pred. No. 18;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                    Sequence 21 BP; 1 A; 10 C; 0 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Microsatellite sequence from clone AGLA248.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rable 7; Page 151; 517pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-FEB-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1992-284684/34.
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                                                                                                                                                                                                                                                 The present sequence represents a phosphorothicate oligonuclectide which
                                                                                             Solid phase synthesis of phosphoro:thioate oligo:nucleotide(s) using new dimeric synthon(s) - useful as anti:sense molecules for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/note= "Phosphorothioate linkages between alternate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleotides (1 and 2, 3 and 4 etc.)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; score 20; DB 18; Length 21; 100.0%; Pred. No. 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 0 A; 10 C; 0 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                 Example 3; Page 20; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 25; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT86583 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 agagagagagagagagag 20
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                                                  WPI; 1997-415290/38.
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Best Local Similarity
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Reese CB;
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Rao MV,
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δλ ВÞ

Query Match

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The sequence is that of a bovine microsatellite sequence obtd. by screening a genomic library of bovine Mbol DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oilgoniceldide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and Mbol sites, the frequency of (Tf) n>9 microsatellites in the bovine genome is estimated at >100. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and downstream of the microsatellite sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amplification of the corresp. microsatellite (using the program obTirghly). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic bovine DNA markers - used in genetic identification,
                                                                                                                                                                                                                                                                                                                                                                                                         PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                were used to generate the required PCR primers for in vitro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437.
                                                                                                                                                                                                                                                                                                                                                   Sequence of microsatellite from clone AGLA209.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23 BP; 12 A; 0 C; 11 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  genetic mapping; traits; amplification; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene mapping, and selective breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Table 7; Page 132; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX00524 standard; mRNA; 24 BP
                                                                                                                                                                                            AAQ33511 standard; DNA; 23 BP
           1 agagagagagagagag 20
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                                          22 AGAGAGAGAGAGAGAG 3
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                                                                                                                                                                                                                                                                                              02-FEB-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9213102-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-AUG-1992,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bos taurus.
                                                                                                                                                                                                                                              AA033511;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          response. The method comprises administering an immunostimulatory monoraded to a non-rodent subject in sufficient quantity to stimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulatory nucleic acid. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rith) for thymidine (T) rich. The netchod is used to vaccinate subjects and/or orthomyxoviridae), bacterial antigens (e.g. herpesylridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. herpesylridae, retroviridae staphylococcus), fungal antigens and/or parasitic antigens. The method is staphylococcus), fungal antigens and/or parasitic antigens. The method is slow useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a finante response and to activate immune cells.

Note: the present sequence may have a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vaccine; cytostatic; virucidal; bactemicidal; fungicidal; anti-parasitic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunostimulatory; tumour; viral infection; bacterial infection;
fungal infection; parasitic infection: cancer; asthma;
infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaccinating against tumors, infectious diseases, allergies and asthmausing immunostimulatory p_{\gamma}\text{-rich} and \tau_{G} nucleic acids -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a method for stimulating an immune
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0
                                                                           100.0%; Score 2); DB 13; Length 22; 100.0%; Pred. No. 18; Cive 0; Mismarches 0; Indels
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Sequence 22 BP; 11 A; 0 C; 10 G; 0 U; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory nucleic acid #819.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krieg AM, Schetter C, Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                    AAF99703 standard; DNA; 22 BP.
                                                                                                                                                                                   1 agagagagagagagag 20
                                                                                                                                                                                                                                   3 agagagagagagagagag 22
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                                                                                                     Best Local Similarity 100.(
Matches 20; Conservative
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25-SEP-1999; 27-SEP-1999;

05-APR-2001.

Synthetic.

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97US-0808474. 97US-0808474. 94US-0320507.

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Gray DM;
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                                                                                                                    03-MAR-1997;
                                                                                                                                                        03-MAR-1997;
                                                                                                                                                                         07-OCT-1994;
                                                 US5856103-A.
                                                                                   05-JAN-1999
               Synthetic.
                                                                                                                                                                                                                                            Clark CL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents a target mRNA for the generation of antisense oligonuclectides (ASO) in a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises:

(a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA.ASO-DNA hybrid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid a series of 13 nearest-neighbour nucleic acid base pair types to provide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Selectively ranking nucleic acid molecules, for inhibitory efficiency - comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence scone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and multiplying
                                                                                                   Target; antisense; selective rank; inhibition; ranking; stability;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Target; antisense; selective rank; inhibition; ranking; stability;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Poly-pyrimidine target sequence for antisense oligonucleotides.
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                                                                  Target sequence #2 for antisense oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               select sequence zones for antisense targeting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 12 A; 0 C; 12 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Pred. ....
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                                30-MAR-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                     Clark CL, Gray DM;
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                                                                                                                      interaction; ss
                                                                                                                                                                                                                                                               03-MAR-1997;
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                                                                                                                                                          Synthetic.
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oligonucleotides (ASO) in a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises:

(a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA.ASO-DNA hybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid determine the fractions into formulas to determine the fractions of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types to provide to produce to nucleic acid base pair types to provide to the nucleic acid antisense sequence; where the resulting ranking to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to
                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents a target mRNA for the generation of antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                             Selectively ranking nucleic acid molecules, for inhibitory efficiency - comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and multiplying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Target; antisense; selective rank; inhibition; ranking; stability;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          select sequence zones for antisense targeting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 BP; 0 A; 12 C; 0 G; 12 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                              Disclosure; Column 13-14; 72pp; English.
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Best Local Similarity
Matches 20; Conserva
WPI; 1999-105098/09.
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WPI; 1999-105098/09.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                targeted to a poly-purine many sequence generated by a method of selectively ranking nucleic acid molecules for inhibitory efficiency.

The method comprises: (a) determining the fraction of each of aset of 13 nearest-neighbour nucleic acid base pair types in a target sequence. Cone RMA-ASO-DNA Mybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid sequence; where the results are ordered to produce a ranking.

The process is used to rank nucleic acid sequences based on the sequence is stability of nucleic acid oligoner binding interactions to select sequence for anticone is an incleic acid oligoner binding interactions to select
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                               Selectively ranking nucleic acid molecules, for inhibitory efficiency - comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide for poly-py::imidine target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 2); DB 20; Length 24; 100.0%; Pred. No. 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                              fractions to determine the fractions and multiplying
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 24 BP; 0 A; 12 C; 0 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Match
Local Similarity 100.0%; Pred. NO. 10,
Local Similarity 0; Mismarches
                                                                                                                                                                                    Disclosure; Column 13-14; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                            sequence zones for antisense targeting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX00527 standard; DNA; 24 BP.
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          03-MAR-1997; 97US-0808474.
                       94US-0320507.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97US-0808474.
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                                                                                           WPI; 1999-105098/09.
                                            (TEXA ) UNIV TEXAS.
                                                                  Clark CL, Gray DM;
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                     07-0CT-1994;
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This oligonucleotide represents an antisense oligonucleotides (ASO) targeted to a poly-pyrimidine mRNA sequence generated by a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises: (a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence room RNA: ASO-DNA hybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid base pair fractions into formulas to nearest-neighbour nucleic acid base pair fractions into formulas to nucleic acid base pair types to provide determined fractions; and concleic acid base pair types to provide determined fractions; and concleic acid base pair types to provide determined fractions; and concleic acid base pair types to provide determined fractions; and concleic acid base pair types to provide determined fractions; and concleic acid base pair types to provide determined fractions; and concleic acid base pair types to provide determined fractions and concleic acid
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RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an antisense
molecule, by heating a triplex to dissociate a Watson-Crick paired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     base pair from a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to select sequence zones for antisense targeting.
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Selectively ranking nucleic acid molecules, for inhibitory efficiency - comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and multiplying
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                                                                                                                                                                                                                                                                                                        Disclosure; Column 13-14; 72pp; English.
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19-JAN-2000; 2000US-0487130.
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The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA:DNA duplex capable of being used as an antisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Watson-Crick pairing. This is useful in antisense therapy, as it controls gene expression by causing transcription to be prevented. The present sequence is an example of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an antisense molecule, by heating a triplex to dissociate a Matson-Crick paired
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Mismatches
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20; Conservative
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                                                               The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA:DNA duplex capable of being used as an antisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Matson Crick pairing. This is useful in antisense therapy, as it controls gene expression by causing transcription to be prevented. The present sequence is an example of a triplex sequence used in the methods of the invention.
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19-JAN-2000; 2000US-0487130.
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          Gray DM,
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    Hoogsteen-paired duplex; Watson-Crick pairing; triplex;
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                                                                          WO200105937-A2.
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                                                                                                                                                               20-JUL-1999;
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                                                                                                       25-JAN-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20;
                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF58000;
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qq ò

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The present invention describes a method for producing a nucleic acid molecule comparising a Hoogsteen-paired RNA:DNA duplace capable of being used as an antisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Watson-Crick pairing. This is useful in antisense therapy, as it controls gene expression by causing transcription to be prevented. The present sequence is an example of a triplex sequence used in the methods of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA:DNA duplex capable of baing used as an antisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Watson-Crick pairing. This is useful in
                                                                                                                                         Generating nucleic acid molecule comprising Hoogsteen-paired RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an antisense molecule, by heating a triplex to dissociate a Watson-Crick paired
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 20; DB 22; Length 24; 100.0%; Pred. No. 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 BP; 12 A; 0 C; 12 G; 0 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid triplex RNA sequence #2.
                                                                                                                                                                                                                                                                                                                              Claim 14; Page 17; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 14; Page 17; 23pp; English.
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Best Local Similarity 100.(
Matches 20; Conservative
Hashem GM;
                                                                    WPI; 2001-159523/16.
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Search completed: August 24, 2002, 22:08:16 Job time: 8849 sec

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(without alignments)
30.766 Million cell updates/sec
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                                                                                                                                     August 24, 2002, 21:57:39; Search time 159.68 Seconds
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/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/ptods_COMB.seq:*
/cgn2_6/ptodata/2/ina/pcomB_seq:*
GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                          383533 seqs, 122816752 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                      OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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20
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	9A-72	JS-08-863-639A-93	4A-8	4A-9	IS-08-808-474A-10	4A-11	4-8	6-1	4-10	4-11	8-19	3-19	3-1	2-1	2B-28	IS-08-267-803B-46	IS-08-222-177A-385	2-5	2-99	3-35	7A-41	3A-2	3A-7	3-13	3-14	3-13	3-14
	JS-08-863-639A-7	3-63	JS-08-808-474A-8	JS-08-808-474A-9	18-47	JS-08-808-474A-1	18-09-235-614-8	JS-09-235-614-9	JS-09-235-614-	JS-09-235-614-1	JS-08-860-038-J	JS-09-580-923-	JS-08-946-138-	JS-09-225-652-	JS-08-469-802B-2	7-80	2-17	IS-08-249-112-5	PCT-US95-06556-	JS-09-580-923-35	IS-08-222-177A-4	JS-09-224-818A-	JS-09-224-818A-	JS-08-860-038-1	JS-08-860-038-	IS-09-580-923-	JS-09-580-923-
	98-86	98-80	08-80	08-80	08-80	08-80	9-23	9 - 23	9-23	9 - 23	98-80	9-28	18-94	19-22	9-46	8-26	18-22	8-24	.0895	9-28	18-22	9-25	9-22	98-86	98-86	928	9-28
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% Query Match	0.001	100.0	0.0	0.001	0.001	100.0	0.0	0.0	0.0	100.0	0.0	0.0	0.0	0.0	0.0	0.0	0.001	0.0	0.0	0.0	0.0	0.0	0.001	0.001	100.0	0.00	0.00.
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C 28 20 100.0 60 1 US-08-222-177A-244 29 20 100.0 78 2 US-08-588-983-24 31 20 100.0 78 2 US-08-588-983-24 C 32 20 100.0 78 2 US-08-588-983-24 C 33 20 100.0 78 2 US-08-588-976-24 C 33 20 100.0 78 2 US-08-588-976-24 C 34 20 100.0 92 1 US-08-222-177A-166 C 34 20 100.0 104 1 US-08-314-596-31 37 20 100.0 104 1 US-08-314-596-31 38 20 100.0 104 3 US-08-314-596-31 39 20 100.0 150 1 US-08-314-596-30 40 20 100.0 150 3 US-08-319-37-31 C 42 20 100.0 150 3 US-08-819-077A-38 C 43 20 100.0 210 1 US-08-222-177A-23 C 43 20 100.0 228 1 US-08-222-177A-43 44 20 100.0 228 1 US-08-222-177A-43 45 20 100.0 243 1 US-07-799-828C-9	RESULT 1 US-08-63-639A-72 US-08-63-639A-72 Sequence 72, Application US/08663639A Patent No. 5981165 GENERAL INFORMATION, Robert S. APPLICANT: Rampal, Jang B. APPLICANT: Caskey, C. T. TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS NUMBER OF SEQUENCES: 95 CORRESEE: Sheldon & Mak STREET: 225 South Lake Avenue, 9th Floor CITY: Pasadena STREET: 225 South Lake Avenue, 9th Floor CITY: Pasadena STREET: 225 South Lake Avenue, 9th Floor CITY: Pasadena STREET: 225 South Lake Avenue, 9th Floor CITY: Dasadena STREET: CACOMPUTER: USA COMPUTER: BEN COMPACHETECT 8 version APPLICATION NUMBER: US/08/863,639A FILING DATE: May 28, 1997 CLASSIFICATION: 435 ATTORNEY AGENT INFORMATION: NAME: JOSSEPH E. Mueth REGISTRATION NUMBER: 20,532 REFERENCY-DOOKET NUMBER: 11859-1 TELECOMUNICATION INFORMATION: NAME: JOSSEPH E. Mueth REGISTRATION FOR SEQ ID NO: 72: SEQUENCE CHARACTERISTICS: LENGTH: 20 DO NO: 72: SEQUENCE CHARACTERISTICS: LENGTH: 20 DO NO: 72: SEQUENCE CHARACTERISTICS: LENGTH: 20 DO NO: 72: STRANDESS: Single TOPPOLOSY: Linear NOLECULE TYPE: Other nucleic acid	Query Match 100.0%; Score 20; DB 2; Length Best Local Similarity 100.0%; Pred. No. 1.4; Matches 20; Conservative 0; Mismatches 0; Indel QY 1 agagagagagagagag 20

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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY:
US-08-808-474A-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Gray, Donald M.
APPLICANT: Clark, Chris L.
TITLE OF INVENTION: METHOD FOR SELECTIVELY RANKING SEQUENCES
TITLE OF INVENTION: FOR ANTISENSE TARGETING
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 20; DB 2; Length 20; 100.0%; Pred. No. 1.4; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: Windows 95 SOFTWARE: Corel Wordberfect 8 version CURRENT APPLICATION DATE: APPLICATION NUMBER: US/08/863,639A FILING DATE: May 28, 1997
                                                                                                                                                                     APPLICANT: Matson, Robert S.
APPLICANT: Coassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
TITLE OF INVENTION: OLIGONUCLEOTIDE REPEAT ARRAYS
NUMBER OF EXQUENCES: 95
                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheldon & Mak
STREET: 225 South Lake Avenue, 9th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: Locke Purnell Rain Harrell
STREET: 2200 Ross Avenue, Suite 2200
                                                                                                       Sequence 93, Application US/08863639A Patent No. 5981185 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: Other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8, Application US/08808474A; Patent No. 5856103; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: Joseph E. Mueth
REGISTRATION NUMBER: 20,532
REFERENCE/DOCKET NUMBER: 11859
TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 796-400
TELEPAX: (626) 795-6321
INFORMATION FOR SEQ ID NO: 93:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 agagagagagagagagag 20
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Matches 20; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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STATE: TexacCOUNTRY: USA
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                    RESULT 2
US-08-863-639A-93/C
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                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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Gaps
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APPLICANT: Clark, Chris L.
TITLE OF INVENTION: METHOD FOR SELECTIVELY RANKING SEQUENCES TITLE OF INVENTION: FOR ANTISENSE TARGETING
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 20; DB 2; Length 24; 100.0%; Pred. No. 1.4; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/808,474A
                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NOBER: 0S/08/808,474A APLICATION NOBER: 03-MAR-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2200 Ross Avenue, Suite 2200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTOKEN, MADEL ADMINES.

RADERRALION NUMBER: 33,732
REPERBENCE, DOCKET NUMBER: UTDAL:001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (214) 740-8000
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                           ALIGNET/ASSATIANCARATION

REGISTRATION NUMBER: 33,732

REPERENCE/DOCKET NUMBER: UTDAL:001

TELECOMMUNICATION INFORMATION:

TELEPHONE: (214) 740-8000

INFORMATION FOR SEQ ID NO: 8:

SEQUENCE CHARACTERISTICS:

LENGTH: 24 base pairs

TYPE: nucleic acid

STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/808,474A FILING DATE: 03-MAR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Best Local Similarity 100.C
Matches 20; Conservative
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STRANDEDNESS: single
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MEDIUM TYPE: Floppy (
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US-08-808-474A-8
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ZIP: 75201-6776
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Dallas
                     Texas
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LENGTH: 24
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APPLICANT: Clark, Chis L.
TITLE OF INVENTION: METHOD FOR SELECTIVELY RANKING SEQUENCES
TITLE OF INVENTION: FOR ANTISENSE TARGETING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Gray, Donald M.
APPLICANT: Gray, Donald M.
APPLICANT: Clark, Chris L.
TITLE OF INVENTION: METHOD FOR SELECTIVELY RANKING SEQUENCES
NUMBER OF SEQUENCES: 37
      100.0%; Score 20; DB 2; Length 24; 100.0%; Pred. No. 1.4; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                          Locke Purnell Rain Harrell
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US-08-808-474A-10/C
; Sequence 10, Application US/08808474A
; Patent No. 5856103
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Mayfield, Denise L.
RECISTRATION NUMBER: 33,732
REFERENCE/DOCKET NUMBER: UTDAI
TELECOMMUNICATION INFORMATION:
TELEFAX: (214) 740-8000
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
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ZIP: 75201-6776
COMBUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
TYPE: Floppy disk
TYPE: TOMPY COMPATIBLE
TOWNSTITER: TOWNST MS-MOSS MS-
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                         Best Local Similarity 100.0
Matches 20, Conservative
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nes 20; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3.
CORRESPONDENCE ADDRESS:
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US-08-808-474A-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                Dallas
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          Query Match
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Patent No. 6165.5...
APPLICANT: CRAY, DONALD M.
APPLICANT: CLARK, CHRISTOPHER L.
TITLE OF INVENTION: AN APPARATUS AND METHOD FOR SELECTIVELY RANKING
TITLE OF INVENTION: AN APPARATUS AND METHOD FOR SELECTIVELY RANKING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
TITLE OF INVENTION: SEQUENCES FOR MAISENSE TARGETING
MANAGEMENT OF TARGETI
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                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/808,474A
                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08/320,507
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (214) 740-8000
TELEFAX: (214) 740-8000
INPORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Mayfield, Denise L. REGISTRATION NUMBER: 33,732 REERENCE/DOCKET NUMBER: UTI TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 08,808
PRIOR FILING DATE: 1997-03-03
PRIOR APPLICATION NUMBER: 08/32C
PRIOR FILING DATE: 1994-10-07
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PAtentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 03-MAR-1997
ATTORNEY/AGENT INFORMATION:
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Matches 20; Conservative
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Matches 20; Conservative
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US-08-808-474A-11
75201-6776
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RESULT 11
US-08-860-038-19/c
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US-09-235-614-11
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                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
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                                                                                                    GREATAL INCOMENTATION:
APPLICANT: GRAY, DONALD M.
APPLICANT: CLARK, CHRISTOPHER L.
TITLE OF INVENTION: AN APPARATUS AND METHOD FOR SELECTIVELY RANKING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
FILE REFERENCE: 91556/66384
CURRENT PELICATION NUMBER: 08/09/235,614
CURRENT FILING DATE: 1999-01-22
PRIOR FILING DATE: 1997-03-03
PRIOR FILING DATE: 1997-03-03
PRIOR FILING DATE: 1994-10-07
NUMBER OF SEQ ID NOS: 38
SCHWARE: PARCENTIN VOF: 2.1
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APPLICANT: CLARK, CHRISTOPHER L.
TITLE OF INVENTION: AN APPRARATUS AND METHOD FOR SELECTIVELY RANKING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
FILE REFERENCE: 91556/66384
CURRENT APPLICATION NUMBER: US/09/235,614
CURRENT FILING DATE: 1999-01-22
PRIOR APPLICATION NUMBER: 08/808,474
PRIOR FILING DATE: 1997-03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: Hybrid DNA US-09-235-614-9
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; Patent No. 6183966
                                             Sequence 9, Application US/09235614
Patent No. 6183966
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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Best Local Similarity 100.09
Matches 20; Conservative
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Best Local Similarity
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                                                                                           GENERAL INFORMATION:
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US-09-235-614-10/c
  RESULT 8
US-09-235-614-9/c
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                                                                       Patent No.
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Gaps
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APPLICANT: CLARK, CHRISTOPHER L.
TITLE OF INVENTION: AN APPARATUS AND METHOD FOR SELECTIVELY RANKING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
FILE REFERENCE: 91556/66384
CURRENT APPLICATION NUMBER: US/09/235,614
CURRENT PILING DATE: 1999-01-22
PRIOR PELING DATE: 1997-03-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence: Hybrid DNA US-09-235-614-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3: Rhone-Poulenc Rorer Inc.
500 Arcola Road, Mailstop 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: ST94090-US TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: FR 94/15162
FILING DATE: 16-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO FR95/01468
FILING DATE: 08-NOV-1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19, Application US/08860038 Patent No. 6287762 GENERAL INFORMATION:
; Sequence 11, Application US/09235614
; Patent No. 6183966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 agagagagagagagagag 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: CROUZET, Joel
APPLICANT: SCHERMAN, Daniel
APPLICANT: WILS, Pierre
                                                                                                                                                                                                                                                                                                                      1994-10-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 agagagagagagagagag 21
                                                                                                                                                                                                                                                                                           PRIOR PALLICATION NUMBER: 08
PRIOR FILING DATE: 1994-10-0
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
                                                   GENERAL INFORMATION:
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STREET:
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APPLICANT: Cameron, Beatrice
APPLICANT: Cameron, Beatrice
APPLICANT: Cameron, Beatrice
APPLICANT: Blanche, Francis
TITLE OF INVEWTION: IMMOBILIZED OLIGON'SCLEOTIDE
FILE REFERENCE: 03804,0138-01
CURRENT APPLICATION NUMBER: US/09/580,923
CURRENT PELLING DATE: 2000-05-26
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 1997-10-08
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
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APPLICANT Glenn Albrecht, Sydney Brenner, David H. Lloyd
TITLE OF INVENTION: Massively Parallel Signature Sequencing by
TITLE OF INVENTION: Ligation of Encoded Adaptors
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Stephen C. Macevicz, Lynx Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                  ;
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                                                                                                                                                                                                                                                                                                           watch 100.0%; Score 20; DB 4; Length 26; Local Similarity 100.0%; Pred. No. 1.4; les 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: oligonucleotide US-09-580-033-19
                                                                                                                                                                                       MOLECULE TYPE: other nucleic acid

DESCRIPTION: /desc = "Oligonucleotide"

US-08-860-038-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 19, Application US/09580923
; Patent No. 6319672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 1, Application US/08946138
; Patent No. 6013445
                                               19:
(610) 454-3816
                                                                                                                                                                                                                                                                                                                                                                                                            l agagagagagagagagag 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                      26 AGAGAGAGAGAGAGAGAG 7
                     TELEFAX: (610) 454-3808
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Crouzet, Joel
Scherman, Daniel
                                                                                           26 base pairs
                                                                                                                   nucleic acid
EDNESS: single
                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; GENERAL INFORMATION:
                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-580-923-19/c
TELEPHONE:
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LENGTH: 26
                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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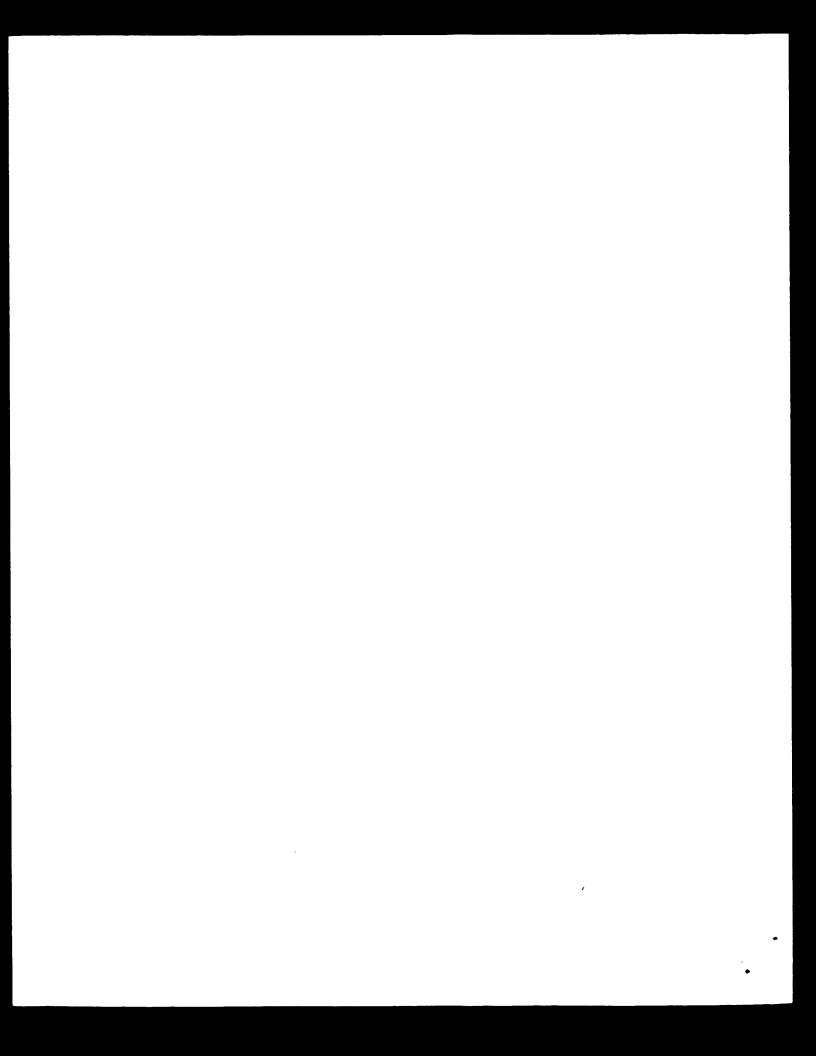
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 20; DB 3; Length 28; 100.0%; Pred. No. 1.4; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM compatible
OPERATING SYSTEM: Windows 3.1/DOS 5.0
SOFTWARE: Microsoft Word for Windows, vers. 2.0
                                                                                                                                                  COMPUTER: IBM compatible OPERATING SYSTEM: Windows 3.1/DOS 5.0 SOFTWARE: Microsoft Word for Windows, vers. 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/946,138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dehlinger & Associates
                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATE:
APPLICATION NUMBER: 08/689,587
FILING DATE: 12-AGG-96
PRIOR APPLICATION DATE:
APPLICATION NUMBER: 08/659,453
FILING DATE: 06-JUN-96
ATTORNEY/AGRAT INFORMATION:
NAME: Stephen C. Macevicz
REGISTRATION NUMBER: 30,285
REFERENCE/DOCKET NUMBER: 808-Lus
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/225,652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              808-1ns
                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/842,608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/09225652 Patent No. 6175002 GENERAL INFORMATION:
3832 Bay Center Place
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (510) 670-9365
TELEFAX: (510)670-9302
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 agagagagagagagagag 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Dehlinger 6
STREET: P.O. Box 60850
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
EDNESS: double
                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 435
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CORRESPONDENCE ADDRESS:
                Hayward
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                USA
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                                                                                 94545
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                                                                COUNTRY:
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0; Gaps
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APPLICANT: Ranum, Laura P.W.
APPLICANT: Chung, Ming-Yi
APPLICANT: Copbi, Huda Y.
TITLE OF INVENTION: Gene Sequence for Spinocerebellar Ataxia
Patent No. 5741645
TITLE OF INVENTION: Type 1 and Method for Diagnosis
NUMBER OF SEQUENCES: 47
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 20; DB 4; Length 28; Best Local Similarity 100.0%; Pred. No. 1.4; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Mueting, Raasch, Gebhardt & Schwappach, P.A.
STREEF: 119 No. 5741645th Fourth Street, Suite 203
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PAPLICATION NUMBER: US/08/469,802B
FILING DATE: 06-JUN-1995
                                            NAME: Vincent M. Powers
REGISTRATION NUMBER: 36,246
REFERENCE/DOCKET NUMBER: 5525-0033
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 324-0880
TELEPHONE: (650) 324-0880
TELEPAX: (650) 324-0880
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 nucleotides
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             110.00030101
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 110
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-305-1217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Mueting, Ann M. REGISTRATION NUMBER: 33,977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 612-305-1225
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 agagagagagagagag 20
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FILING DATE: 15-APR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 32 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: DNA
US-08-469-802B-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
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100.0%; Score 20; DB 1; Length 32;

Query Match



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August 24, 2002, 21:19:22; Search time 4810.92 Seconds (without alignments) 56:110 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                  27472414
GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                      13736207 seqs, 6748477542 residues
                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                          OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                   1 agagagagagagagag 20
                                                                                                                                                                                                                                                                                         IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                      Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                       US-10-077-383-29
                                                                                                                                                                                                                                                                                         Scoring table:
                                                                                                                                                                                                                              Perfect score:
                                                                                                                                                                                                                                                                                                                                                      Searched:
                                                                                                                                                                                                                                                   Sequence:
                                                                                                                        Run on:
                                                                                                                                                                                                          Title:
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1: em_estba:*
2: em_esthum:*
3: em_esthum:*
4: em_esthu:*
5: em_esto:*
6: em_estpl:*
7: em_estro:*
8: em_htc:*
10: gb_est2:*
11: gb_htc:*
12: gb_gs:*
13: em_gss_hum:*
14: em_gss_lnv:*
15: em_gss_lnv:*
16: em_gss_lv:*

ESI: *

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description			 	AZ627978 1M0476L04	AZ633751 1M0489111	AZ822888 2M0096G06	AZ389918 1M0151E05	AZ510124 1M0354K22	AZ659095 1M0536E18	AZ664804 1M0545H24	AZ342914 1M0076C22	AZ579594 1M0367010	AZ641486 1M0504J06	AZ666145 1M0548C02	AZ771239 1M0573F15	AZ803946 2M0064P06	AZ813253 2M0080M20
SUMMARIES	ID	AZ433566	AZ394897	AZ589098	AZ627978	AZ633751	AZ822888	AZ389918	: A2510124	: AZ659095	~	_	: AZ579594	AZ641486	: AZ666145	: AZ771239	: AZ803946	AZ813253
	B0 1	12	. 12	7 .	7.5	12	12	12	12	12	12	•	•		12	12	12	12
	Query Match Length DB	20	5.2	7 6	7	22	23	25	25	25	25	56	26	56	26	56	56	56
ф	Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
	Score	20	202	0 0	07	20	20	20	20	20	20	20	20	20	20	20	20	50
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	Result No.	O	ပ		υ	U		υ	ပ		U	υ		U			U	O

/strain="C57BL/6J" /db_xref="taxon:10000" /clone="UGCLM0219C02" /clone_lib="Mouse 10kb plasmid UUGCIM library"

Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends High quality sequence stop: 20. Location/Qualifiers

/organism="Mus musculus"

source

FEATURES

20 100.0 26 12 AZ632991 AZ632991 IMC487M21 20 100.0 27 12 AZ655331 AZ655391 IMC487M21 20 100.0 27 12 AZ76549 AZ462349 AZ462349 20 100.0 27 12 AZ655331 AZ462249 AZ462249 AZ462249 20 100.0 28 12 AZ462249 <	ALIGNMENTS ALIGNMENTS ALIGNMENTS N 19021902R Mouse 10kb plasmid UUGCLM library Mus musculus genomic clone UUGCLM0219022 R, DNA sequence. A2433566 A2433566. GI:10537579 GSS. House mouse. Musculus musculus ENATYOTA: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Manmalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus. 1 (bases 1 to 20) Losses 1 to 20) Losses 1 to 20) Lonn, D. Aoyaqi.A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Tingey, A., von Niederhausern, A. Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts University of Utah Genome Center Utah Genome Center Utah Genome Center Utah Genome Center Utah Genome C
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	RESULT 1 A2413566/C LOCUS DEFINITION ACCESSION VERSION VERSION CENTER ORGANISM TITLE JOURNAL COMMENT

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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                             10.5 Kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114|gblAF129072.1), a copy.number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified the sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically.competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance.
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                                                  /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv: Purified genomic DNA from M.
musculus C57EL/60 (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
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Dunn, D., Aoyaqil,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Meiss,R.
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1M0397819R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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Seq primer: CACACAGGAAACAGCTATGACC
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University of Utah Genome Center
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Location/Qualifiers
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Fax: 801 585 7177
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                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a was hydrodynamically sheared by repeated passage through a was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarcse gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored mouse DNA was annealed to
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                         /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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Insert Length: 10000 Std Error: 0.00
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Location/Qualifiers
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/clone="UUGC1M0158H11"
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Fax: 801 585 7177
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Fax: 801
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                                                                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electropheresis. Vector DNA was prepared from a derivative of pWDA2 (qi[4732114 [qb] AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically competent E. coll XL10-Gold (Stratagene) cells and solected for ampicillin resistance."
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Rellly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Welss,R.
                                                                                                   /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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             /clone_lib-"Mouse 10kb plasmid UUGClM library"
/sex-"Male"
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/clone="UUGC1M0397B19"
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/strain="C57BL/6J"

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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarcose gel electrophoresis. Vector DNA was prepared from a derivative of pwBV2 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of phasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-completent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
, M., Rose, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
                                                                                                  /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
hote="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/clone_"UUGCIM0476L04"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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Insert Length: 10000 Std Error: 0.00
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/organism="Mus musculus"

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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/comments/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam, H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,
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2M0096G06R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0096G06 R, DNA sequence.
                                                                                                                                                                                                                                          /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC1M library"
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Plate: 0006 row: G column: 06
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah Genome Center
University of Utah
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Location/Qualifiers
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/clone="UUGC1M0489111"
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/strain="C57BL/6J"
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Fax: 801 585 7177
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Ladoracupy mouse DNA Resources, documents, dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pw042 (gil432114) glbAF129072.11, a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and solected for ampicillin resistance."
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                    /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"
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                                                                                      /clone_lib="Mouse 10kb plasmid UUGC1M library"
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Class: plasmid ends
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                              /db_xref="taxon:10090"
                                                        /clone="UUGC2M0096G06"
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Insert Length: 10000 Std Erro
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Matches 20; Conservative
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Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polymerlectiede kinase. Adaptor oligonuclecticates were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 25)
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
hote="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                          /clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                    /organism="Mus musculus"
/strain="C57BL/6J"
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Class: plasmid ends
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Contact: Robert B. Weiss
University of Utah Genome Center
                                                                                                                            /db_xref="taxon:10090"
/clone="UUGC1M0354K22"
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Insert Length: 10000 Std Erro:
Plate: 0536 row: E column: 18
         Location/Qualifiers
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                                                                                                                                                                                                                       /sex="Male"
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                   http://www.jar.cogresources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclactide kinase. Adaptor oligonuclactides were ligated to the blunt ends in high molar axess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114]sb[AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically competent E. coll XL10-Gold (Stratagene) cells and selected for ampizillin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 25)
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1M0354K22F Mouse 10kb plasmid UUGClM library Mus musculus genomic
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                                                                                                                                                                                                                                    /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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                                                       /strain="C57BL/6J"
/db_xref="taxon:10000"
/clone="UGCIMO151E05"
/clone_lib="Mouse 10kb plasmid UGGCIM library"
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                          /organism="Mus musculus"
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Seq primer: CGTTGTAAAACGACGCCAGT
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University of Utah Genome Center
University of Utah
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Insert Length: 10000 Std Erro
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Fax: 801 585 7177
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                               /organism="Mus musculus"
/strain="C57BL/65"
/db_xref="kacon:10090"
/clone="UUGCIM0356E18"
/clone="UUGCIM0356E18"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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Insert Lenghh: 1000 Std Error: 0.00
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Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (g114732114]pb[AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Contact: Robert B. Weiss
Contact: Robert Und Genome Center
University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Dunn, D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Relly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                                                                                                                                                                                                                                                /lab host="E. Coli strain XLIO-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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Insert Length: 10000 Std Error: 0.00
Plate: 0076 row: C column: 22
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/clone="UUGC1M0545H24"
High quality sequence stop: 25.
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Email: ddunn@genetics.utah.edu
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Musculus C57BL/60 (male) was obtained from the Jackson
Laboratory Mouse DNA Resource from the Jackson
Laboratory Mouse DNA Resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD4 (gil4732114|pblAR129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
chemically-competent E. coll XL10-Gold (Stratagene) cells
and selected for ampizillin resistance."
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Bukaryota; Metazoa; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 26)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
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/clone="UUGCLM0076C22"
/clone_lib="Mouse 10kb plasmid UUGCLM library"
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Insert Length: 10000 Std Error: 0.00
Seq primer: CGTTGTAAAACGACGGCCAGT
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Context: Robert B. Weiss
University of Utah Genome Centar
University of Utah
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                                                                              Location/Qualifiers
                                                                                                                                                          /strain="C57BL/6J
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Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWN42 (gile/32114 [gb] ART29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected completion mapping and selected in mapicillin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Dunn, D., Aoygal, M., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly

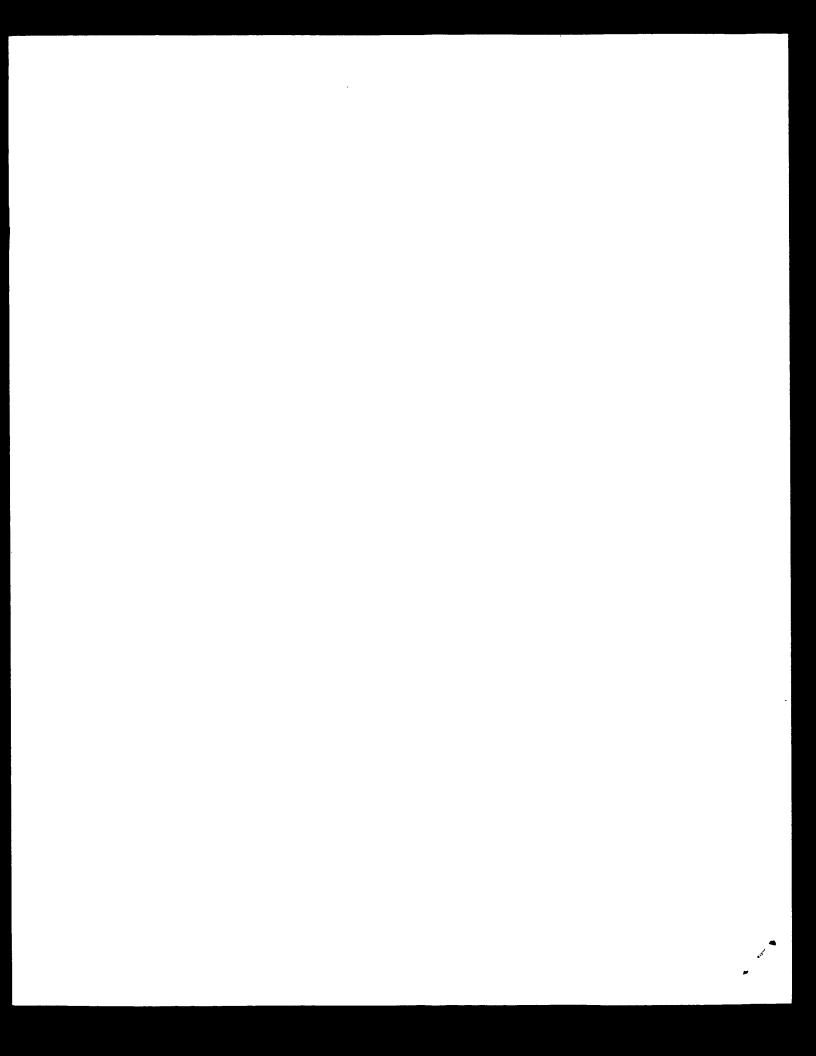
M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.,
and Wright, D., Weiss, R.
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hote="Vector: PWD42nv; Purified genomic DNA from M.
musculus G57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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Seq primer: CGTTGTAAAACGACGCCAGT
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University of Utah Genome Center
University of Utah
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/clone="UUGC1M0367010"
                                                                                                  High quality sequence stop: 26.
Location/Qualifiers
                                                                                                                                                                                                                                           /strain="C57BL/6J
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                                                                    Class: plasmid ends
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 26)
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1M0548C02F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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                      Plate: 0504 row: J column: 06
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University of Utah Genome Center
University of Utah
                                                                                                                                                                                                         /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0504J06"
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Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymerlectiede kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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26 bp DNA linear GSS 16-FEB-200
1M0573F15F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0573F15 F, DNA sequence.
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Mouse whole genome scaffolding with paired end reads from 10kb
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Email: ddunn@genetics.utah.edu
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Contact: Robert B. Weiss
University of Utah Genome Center
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/clone="UUGC1M0548C02"
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Location/Qualifiers
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Search completed: August 24, 2002, 21:19:25 Job time: 20528 sec



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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Pizza,M., Staria and Statement (17): 2.20-MAY-1999;
PIZZA MARIAGRAZIA (IT): SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT): GRANDI GIIDO (IT); MASIGNANI VEGA (IT)
Location/Qualifiers
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Neisserial antigens
Patent: W 9924578-A 45 20-MAY-1999;
PIZZA MARIAGRAZIA (IT): SCARLATO VINCENZO (IT): RAPPUOLI RINO (IT);
CHIRON SPA (IT): GRANDI GUIDO (IT): MASIGNANI VEGA (IT)
                                                                                                                                                                                                                                    Pizca,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V. Neisserial antigens
Patent: w 0.924578-A 35 20-MAY-1999;
Pizca MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Neisserial antigens
Patent: WO 9924578-A 25 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Neisserial antigens
Patent: WO 9924578-A 35 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Sequence 25 from Patent WO9924578.
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Sequence 35 from Patent W09924578.
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/db_xref="taxon:32644"
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
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Sequence 73 from Patent WO9924578.
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/db_xref="taxon:32644"
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                                                                                                              Neisserial antigens
Patent: WO 9924578-A 45 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Patent: WO 924578-A 55 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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1 (bases 1 to 1)
Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
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Sequence 55 from Patent WO9924578.
A96022.
A96022.1 GI:6779903
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/db_xref="taxon:32644"
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A96012
A96012.1 GI:6779898
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Neisserial antigens
Patent: WO 9924578-A 55 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Meisserial antigens
Patent: WO 9234578-A 73 20-MAY-1999;
PIZZA MARIAGRAZIA (IT): SCARLATO VINCENZO (IT): RAPPUOLI RINO (IT):
CHIRON SPA (IT): GRANDI GUIDO (IT): MASIGNANI VEGA (IT)
Location/Qualifiers
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Patent: WO 9924578-A 73 20-MAY-1999;
PIZZA MAKIAGRAZIA (IT): SCARLATO VINCENZO (IT): RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
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Sequence 109 from Patent WO9924578.
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/db_xref="taxon:32644"
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Pizza,Estal antigens
Patent: WO 9224578-A 91 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Neisserial antigens
Patent: WO 924278-A 91 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Pizza,M., 9924578-A 109 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Patent: WO 9924578-A 109 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza.M. Scarlato.V., Rappuoll,R., Grandi,G. and Masignani,V.
Pizza.M. Sozrlato.Sozrlato.V.
Pizza.M. Scarlato.V.
CHIRON SPA (IT): GRANDI GUIDO (IT): MASIGNANI VEGA (IT)
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Description	Human mutant KCNO3	Human mutant KCNO3	Segmence encoding	Seguence encoding	Plasmid DEZZ318 +h	Plasmid pEzz318 th	CAPL trinucleotide	CAPL trinucleotide	Human TUB gene pro
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KCNQ2;	KCNQ3; hu	man;	ne;	tassium	el; diagnosis; prognosis;
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WPI;	1999-312	2938/26.			

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proteins KCNO3 and KCNO3. Detecting mutations in sequences that encode KCNO2 or KCNO3. or the loss of one copy of these genes, is used for diagnosis and prognosis of benign familial neonatal epilepsy (BFNE). Juvenile myotonic epilepsy (JANE) or rolandic epilepsy (RFN). Cells (or transgenic animals) that express wild-type or mutant KCNO2 or 3 (also the proteins themselves in cell-free form) are used to screen for agents that can be used to treat or prevent these forms of epilepsy. Fragments of the encoding mutations or for isolation of related sequences, while the complete sequences may be used in gene therapy to provide wild-type protein. Antibodies specific for mutant or wild-type proteins are used in carional design of drugs and therapeutically (in replacement therapies). The forms of epilepsy associated with mutations in KCNQ2 and 3 sequences can now be diagnosed early (before symptoms are manifest), and better treatment options will be available.

AAX57074-X57139 are primers used in the method of the invention.
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Nucleic acid encoding potassium channels KCNQ2 and 3
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KCNQ2 or KCNQ3, or the loss of one copy of these genes, is used for diagnosis and prognosis of benign familial neonatal epilepsy (BRNE), juvenile myctoric epilepsy (BRNE) or rolandic epilepsy (RR). Cells (or transgenic animals) that express wind-type or mutant KCNQ2 or 3 (also the proteins themselves in cell-free form) are used to screen for agents that can be used to treat or prevent these forms of epilepsy. Fragments of the encoding nucleic acids are used as probes or primers, either for detecting mutations or for isolation of related sequences, while the complete sequences may be used in gene therapy to provide wild-type protein. Antibodies specific for mutant or wild-type proteins are used as diagnostic reagents and for drug screening. The KCNQ2 and 3 proteins are useful in rational design of drugs and therapeutically (in
                                                                                                                                                                                                                 replacement therapies). The forms of epilepsy associated with mutations in KNO2 and 3 sequences can now be diagnosed early (before symptoms are manifest), and better treatment options will be available.

AAX57074-X57139 are primers used in the method of the invention.
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                            A DNA segment with a nucleotide sequence from about 130 to about 918 of the sequence given here is claimed. Also claimed are antibodies which immunoreact with hurFH and the claimed peptides (given in AAP80713). The antibodies may be used in immunoassays for detection of hurFH. The claimed peptides may be used to inhibit the binding of hurF to coagulation factor VII/VIIa in vivo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    10-SEP-1990 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; ; pp; English.
Disclosure; ; pp; English.
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                                                                                                                                                               Gaps
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                                                                                                                                                        0; Indels
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                                                                                    / Match 0.0%; Score 0; DB 9; Length 3; Local Similarity 100.0%; Pred. No. 0; nes 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OMTKY3; turkey ovomucoid third domain peptide inhibitor;
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Sequence 3 BP; 0 A; 0 C; 0 G; 0 T; 3 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ85491 standard; cDNA; 3 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmid pEZZ318.thy2.
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Claim 2; Page 56; 70pp; English.
96WO-US02108.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            osteosarcoma; therapy; ss.
                                                    (HYBR-) HYBRIDON INC
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                                                                                                                                      WPI; 1996-393400/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAPL trinucleotide.
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16-FEB-1996;
                          17-FEB-1995;
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Von Hofe E;
                                                                                                         Von Hofe E;
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                                                                                                                                                                                                                                                                                                                                                                                      Protein inhibitors of serine proteinase(s), e.g. furin, deriv. from turkey ovomucoid third domain - used as laboratory reagents to study the proteinase(s), or as chemotherapeutic agents to treat diseases associated with them
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                                                                                          OMTKY3; turkey ovomucoid third domain peptide inhibitor;
furin inhibitor; protease inhibitor; pEZ2318.thy2; probe;
Escherichia coli; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3 BP; 0 A; 0 C; 0 G; 0 T; 3 other;
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                                         18-AUG-1995 (first entry)
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                                                                                                                                                                                                                                                                                                                                   Anderson S, Laskowski M;
                                                                      Plasmid pEZZ318.thy2.
                                                                                                                                                                                                                                                                                                        (LASK/) LASKOWSKI M.
                                                                                                                                                                                                                                                                                                                                                           WPI; 1995-066900/09.
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                                                                                                                                                                                                                                                                                            (ANDE/) ANDERSON S.
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                                                                                                                                                      Synthetic.
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RESULT

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Gaps
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                                                                                                                                                                                     Agrawal S, Engebraaten O, Fodstad O, Hovig E, Maelandsmo GJ;
                                                                                                                                                                                                                                                                                                                                                                           Synthetic oligo:nucleotide(s) inhibiting CAPL gene expression -
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                                                                                                                                                                                                                                                                                                                                                                                                                        useful to inhibit metastatic cancer, partic. osteosarcoma
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                                                                                                               (NORA-) NORWEGIAN RADIUM HOSPITAL RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HYBR-) HYBRIDON INC.
(NORA-) NORWEGIAN RADIUM HOSPITAL RES FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 56; 70pp; English.
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95US-0391375.
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Sequence 3 BP; 0 A; 0 C; 3 G; 0 U; 0 other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
Novel antisense oligonucleotides capable of inhibiting CAPL gene expression may include the trinucleotide GUC (AAT33326, given in 5' to 3' direction) found in codon 14 of CAPL mRNA. These and other antisense oligonucleotides (AAT33327-36) complementary to specific regions of the CAPL gene (see also AAT33346), as well as CAPL-specific ribozymes (AAT3337-40) can be administered to a patient as a means of inhibiting metastatic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; TULP; neurosensory defect; ret.ina; retinal dystrophy; probe;
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                                                                                                                                                                                                                                                                                                                                                                                       Score 0; DB 17; Length 3;
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                                                                                                                                                                                                                                                                                                                                              Query Match 0.0%; Score 0; DB 1,
Best Local Similarity 0.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                  Sequence 3 BP; 0 A; 1 C; 1 G; 1 U; 0 other;
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30-APR-1997;
01-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalian proteins expressed in retina and brain, useful for producing antibodies and for diagnosing neurosensory defects including cochlear degeneration, peripheral retinal degeneration and cone-rod retinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for the treatment of neurosensory degenerative conditions e.g. retinal dystrophies. The present sequence is a probe used to isolate the novel genes of the present invention.
                                            Gaps
                                                                                                                                                                                                                                                                                                    Human; TULP; neurosensory defect; retina; retinal dystrophy; probe;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to human and murine cDNAs from a
                                         1; Indels
           Score 0; DB 21; Length 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      North M, Nishina P, Noben-Trauth K, Naggert J;
                        Pred. No. 0;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Columns 81-82; 61pp; English.
                                                                                                                                                                           AAA94655 standard; DNA; 3 BP.
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97US-0904699.
97US-0932306.
           0.08;
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Query Match
Best Local Similarity 0.0%;
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                                                                                                                                                                                                                                      15-JAN-2001 (first entry)
                                                                                                                                                                                                                                                                   Human TUB gene probe #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AXYS-) AXYS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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01-AUG-1997;
17-SEP-1997;
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10-APR-1996;
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WO9500528-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to human breast cancer expressed polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AALO7544-AAL26789) and methods of assessing whether a patient is afflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells.
          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New peptide useful as a marker for the diagnosis of breast cancer
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                                                                                                                                                                                                     Human breast cancer expressed polynucleotide 12701.
                                                                                                                                                                                                                             Human; breast cancer; cell marker; cytostatic; ss.
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Pred. No. 0;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lillie J, Xu Y, Wang Y, Steinmann K;
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                                                                                                                          AAL20244 standard; cDNA; 3 BP.
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15-WAY-2000; 2000US-0205230.
09-JUN-2000; 2000US-0211315.
25-JUL-2000; 2000US-0220534.
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            Conservative
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Best Local Similarity
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The invention relates to human breast cancer expressed polynucleotides (AAL07544-AAL26789) and methods of assessing whether a patient is safflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells. The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic
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                                                        Human breast cancer expressed polynucleotide 12701.
                                                                                                                     Human; breast cancer; cell marker; cytostatic; ss.
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         bFGF binding oligomer core sequence #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lillie J, Xu Y, Wang Y, Steinmann K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 2245; 3695pp; English.
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09-JUN-2000; 2000US-0211315.
25-JUL-2000; 2000US-0220534.
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07-DEC-2001 (first entry)
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Best Local Similarity
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                                                                                                                                                                                       Homo sapiens.
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AAQ81665
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tumours and to reduce the risk of metastasis. They can be used as diagnostic reagents to determine the presence of thrombin, or used in
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                                                                                                                                                                                                                                                                                                                                                                                                      New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
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0.0%; Pred. No. 0;
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                                                                                    94WO-US06884.
                                                                                                                                             93US-0079677.
94US-0179491.
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                                                                                                                                                                                                                                  (PHAR-) PHARMAGENICS INC.
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                                                                                                                                                                                                                                                                                           Beutel BA, Joesten ME;
                                                                                                                                                                                                                                                                                                                                                WPI; 1995-051992/07.
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Best Local Similarity
                                                                                                                                          18-JUN-1993;
07-JAN-1994;
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07-JAN-1994;
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The sequences given in AAQ81642-95 are oligonucleotides which medulate the activity of basic fibroblast growth factor (bFGF) by binding to the bFGF protein. Most easp. the sequences given in AAQ81664-67 represent core sequences of which at least one is present in each of the binding oligos. These oligonucleotides may form a single strand, double strand, a stem-loop structure, a bubble structure, pseudoknot or a closed, circular structure. bFGF binds to high affinity receptor and low affinity hearing it was not the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumours and to reduce the risk of metastasis. They can be used as diagnostic reagents to determine the presence of thrombin, or used in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell surface. These oligonuclectides bind to bFGF in competition with its receptor and heparin. These oligonuclectides may inhibit or enhance the activity of bFGF. Particularly, they inhibit neovascularisation so they can be used to suppress growth of solid tumours and to reduce the risk of metastasis. They can be used as
                                                                                                                                                                                                                                                      factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Basic; fibroblast growth factor; bFGF; stem-loop structure; bubble structure; pseudoknot; receptor; heparin; competition; inhibition; enhance; neovascularisation; solid tumour; cancer; metastasis; diagnosis; gene therapy; ss.
                                                                                                                                                                                                                  New oligo-nucleotide(s) that bind to basic fibroblast growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.0%; Score 0; DB 16; Length 4; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 25; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ81665 standard; RNA; 4 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94WO-US06884.
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(PHAR-) PHARMAGENICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHAR-) PHARMAGENICS INC.
                                                                   Beutel BA, Joesten ME;
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                                                                                                                                           WPI; 1995-051992/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy.
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The sequences given in AAQ81642-95 are oligonucleotides which modulate the activity of basic fibroblast growth factor (bFGF) by binding to the bEGF protein. Most esp. the sequences given in AAQ81664-67 represent core sequences of which at least one is present in each of the binding oligos. These oligonucleotides may form a single strand, double strand, a stem-loop structure, a bubble structure, a pseudoknot or a closed, circular structure. BrGF binds to high a ffinity receptor and low affinity heparin-like molecules on the circular structure or large oligonucleotides bind to bFGF in competition with its receptor and heparin. These oligonucleotides may inhibit or enhance the activity of DFGF. Particularly, they inhibit or enhance the activity of DFGF. Particularly, they inhibit necessis and to reduce the risk of metastasis. They can be used as diagnostic reagents to determine the presence of thrombin, or used in
New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
                                                                                                              Claim 3; Page 25; 44pp; English.
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Search completed: August 24, 2002, 22:08:16 Job time: 8849 sec

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Query Match 0.0%; Score 0; DB 16; Length 4; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels

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Sequence 4 BP; 0 A; 2 C; 1 G; 1 U; 0 other;

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Sequence 62, A sequence 7, A sequence 7, A sequence 22, B sequence 22, A sequence 2, A sequence 62, A sequence 62, A sequence 62, A sequence 2, A sequence 36, A

US-07-791-213D-62 US-08-268-679B-7 US-08-295-743-22 US-08-295-743-22 US-08-295-743-22 US-08-295-743-22 US-08-602-036A-2 US-08-602-036A-2 US-08-602-036A-2 US-08-293-150A-62 US-08-293-150A-62 US-08-502-314A-2 US-08-502-314A-2 US-08-612-407A-2 US-08-612-407A-2

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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
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/cgn2_6/ptodata/2/ina/ROTOS_COMB.seq:*
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       383533 seqs, 122816752 residues
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Listing first 45 summaries
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Description	Sequence 702, App	702,	8, Ap	8	16,	16,	16,	16,	28,	28,	35,	35,	35,	35,	35,	35,	3,	'n	15,	15,	16,	16,	16,	16,	46,	46,	62,
ID	PCT-US93-00977-702	PCT-US93-00977-702	US-08-268-679B-8	US-08-268-679B-8	US-08-457-274A-16	US-08-457-274A-16	US-08-484-192-16	US-08-484-192-16	US-08-726-464B-28	US-08-726-464B-28	-00-016-	US-09-016-520-35	US-09-130-973-35	US-09-130-973-35	US-09-477-902-35	US-09-477-902-35	US-08-361-024-3	US-08-361-024-3	US-09-411-862A-15	Н	US-09-411-862A-16	US-09-411-862A-16	ç	PCT-US95-35758-16	US-07-791-213D-46	US-07-791-213D-46	US-07-791-213D-62
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SUMMARIES

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ALIGNMENTS	PCT-US93-00977-702 Sequence 702, Application PC/TUS9300977 Sequence 702, Application PC/TUS9300977 GENERAL INFORMATION: TITLE OF INVENTION: MUDHERS PER SEQUENCES: 711 CORRESPONDENCE ADDRESS: ADDRESSEE: Knobbe, Martens, Olson, and Bear STRET: 620 Newport Center Dr. Sixteenth Floor CITY: Newport Beach STATE: CA COMPTRY: USA COMPTRY: USA COMPTRY: Bach STATE: Ploppy disk COMPTRY: Bach COMPTRE REDABLE FORM: MEDIUM TYPE: Ploppy disk COMPTRE: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 COMPTRE: IBM PC COMPATION: MEDIUM TYPE: Ploppy disk COMPTRE: 19930129 CLASSIFICATION NUMBER: 34,115 REFERENCE/DOCKET NUMBER: #117ACHI.006H TELECHONE: 714-760-9502 INFORMATION FOR SEQ ID NO: 702: SEQUENCE CHARACTERISTICS: LEWITH: 1 TYPE: NUCLEIC ACID STRANDEDNES: double TOPOLOGY: linear MULECULE TYPE: NO HYPOTHETICAL: NO MATI-SENSE: NO ANTI-SENSE: NO ANTI-SENSE: NO PCT-USSI-OU977-702	Query Match 0.0%; Score 0; DB 5; Length 1; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels	Qy 1 n 1	Db 1 A 1

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STRANDEDNESS:
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                                                                         TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
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                                                                                                                      ADDRESSEE: Knobbe, Martens, Olson, and Bear
STREET: 620 Newport Center Dr. Sixteenth Floor
CITY: Newport Beach
                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: WIMMER, ECKARD; MOLLA,
APPLICANT: AKHTERUZAMAN; PAUL, ANIKO V.
TITLE OF INVENTION: DE NOVO CELL-FREE
TITLE OF INVENTION: SYNTHESIS PICORNAVIRUS
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY AGENT INFORMATION:
NAME: Altman, Daniel E.
REGISTRATION NUMBER: 34,115
REFERNCE/DOCKET NUMBER: HITACHI.006H
TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2: MORGAN & FINNEGAN, L.L.P. 345 PARK AVE.
                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US93/00977 FILING DATE: 19930129
                                 Sequence 702, Application PC/TUS9300977 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORD PERFECT # 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 702:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: cDNA to mRNA HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 double
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MEDIUM TYPE: FLOPPY I
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                                                                                              NUMBER OF SEQUENCES: 7. CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEIC ACID
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                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ANTI-SENSE: NO PCT-US93-00977-702
RESULT 2
PCT-US93-00977-702/c
                                                                                                                                                                                                                USA
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                                                                                                                                                                                                CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-268-679B-8
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Query Match 0.0%; Score 0; DB 1; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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APPLICANT: AKHTENUZAMAN; PAUL, ANIKO V.
TITLE OF INVENTION: DE NOVO CELL-FREE
TITLE OF INVENTION: SYNTHESIS PICORNAVIRUS
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                  0887-4095 US2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MORGAN & FINNEGAN, L.L.P.
APPLICATION NUMBER: US/08/268,679B FILING DATE: 30-JUN-1994
PRIOR APPLICATION DATA: APPLICATION NUMBER: 07\846,914
FILING DATE: 06-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORD PERFECT # 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/268,679B
FILLING DATE: 30-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    435
mRR: 07\719,761
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FRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                          APPLICATION NUMBER: 07\719,761
FILING DATE: 24-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: MARIA C.H. LIN
RESISTRATION NUMBER: 29,333
REFERENCE/DOCKET NUMBER: 0897-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8, Application US/08268679B Patent No. 5674729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DESCRIPTION: OLIGONUCLEOTIDE
                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                            TELEX: 421792
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 07\84
FILING DATE: 06-MAR-1992
CLASSIFICATION: 435
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FILING DATE: 24-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: MARIA C.H. LIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORIGINAL SOURCE: N.A.
; POSITION IN GENOME: N.A.
US-08-268-679B-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 345 PARK AVE. CITY: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                        CLASSIFICATION:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                              US-08-457-274A-16/C
Sequence 16, Application US/08457274A
Patent No. 57340B
GENERAL INFORMATION:
APPLICANT: Scott, Jeffrey G.
APPLICANT: TARBANI
TITLE OF INFERTION:
WUMBER OF SEQUENCES: 29
                                                                                                                                                                                                        1; Indels
                                                                                                                                                         Query Match 0.0%; Score 0; DB 1; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIDLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PactentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/457,274A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEE: Nixon, Hargrave, Devans & Doyle
: P.O. Box 1051, Clinton Square
Rochester
               STRAIN: Learn-PyR
DEVELORWHIALS, STRGE: Adult
POSITION IN GENOME:
CHROMOSOME/SEGMENT: Chromosome 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CHROMOSOME/SEGMENT: Chromosome 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: GOLDMan, Michael L.
REGISTRATION NUMBER: 30,727
REFERNCE/POCKET WUMBER: 1960;
TELECOMMUNICATION INFORMATION:
TELEPHONE: 716-263-1304
TELEFAX: 716-263-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DEVELOPMENTAL STAGE: Adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 POSITION IN GENOME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New York
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ORGANISM: MUS
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                                                                                               US-08-457-274A-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Scott, Jeffrey G. APPLICANT: Scott, Jeffrey G. APPLICANT: Tomita, Takashi TITLE OF INVENTION: Cytochrome P450lpr Gene and Its Uses CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.0%; Score 0; DB 1; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/457,274A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Nixon, Hargrave, Devans & Doyle
STREET: P.O. Box 1051, Clinton Square
CITY: Rochester
| REGISTRATION NUMBER: 29,323 | REFERENCE/DOCKET NUMBER: 0887-4095 US2 | TELECOMMUNICATION INFORMATION: | TELEFHONE: (212) 758-480 | TELEFAX: (212) 751-6849 | TELEFAX: (212) 751-6849 | TELEFAX: (212) 751-6849 | INFORMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 16, Application US/08457274A; Patent No. 5734086
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: GOldman, Michael L.
REGISTRATION NUMBER: 30,727
REFERENCE/DOCKET NUMBER: 19603
TELECOMMUNICATION INFORMATION:
TELEPHONE: 716-263-1304
TELEPENCE: 716-263-1304
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                         DESCRIPTION: OLIGONUCLEOTIDE POTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Musca domestica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Floppy disk
                                                                                                                                                                                                                                                                                                                                     ORIGINAL SOURCE: N.A.
POSITION IN GENOME: N.A.
US-08-268-679B-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 2 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                      SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                             NUCLEIC ACID
                                                                                                                                                                                                               STRANDEDNESS: SING
TOPOLOGY: UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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                                                                                                                                                                                                                                                         MOLECULE TYPE:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/934,387
FILING DATE: 21-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/484,192
                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 28, Application US/08726464B Patent No. 5932546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: BARRETT, Ronald W. APPLICANT: DOWER, William J. APPLICANT: CWIRLA, Steven A. APPLICANT: JOHNSON, Sherril S. APPLICANT: WRIGHTON, Nicholas C. APPLICANT: DUFFIN, David J.
                                                                                                                                                          E: MORRISON & FOERSTER 755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-813-5600
TELEFAX: 415-494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28,216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 706141
INFORMATION FOR SEQ ID NO: 16:
                   LEUNG, LAWRENCE
                                                                                                                                                                                                                                                                                                      Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 2 base pairs
TYPE: nucleic acid
                                       VERMAAS, ERIC
TOOLE, JOHN J
                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: GRACEY, NANCY J. REGISTRATION NUMBER: 2
APPLICANT: LATHAM, JOHN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                   STREET: 755 ALCOUNTY: PALO ALTO
                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                            94304
                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-726-464B-28
                                     APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-484-192-16
                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                    APPLICANT: ALBRECHT, GLENN
APPLICANT: ALBRECHT, GLENN
APPLICANT: LATHAM, JOHN
APPLICANT: LEDNG, LAMENCE
APPLICANT: VERMAAS, ERIC
APPLICANT: TOOLE, JOHN J.
APPLICANT: TOOLE, JOHN J.
TITLE OF INVENTION: APTAMERS SPECIFIC FOR BIOMOLECULES AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 21-AUG-1992
ATTORNEY/AGENT INFORMATION:
NAME: GRACEY, NANCY J.
REGISTRATION NUMBER: 28.216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/484,192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: MORRISON & FOERSTER STREET: 755 PAGE MILL ROAD
                                                                            Sequence 16, Application US/08484192; Patent No. 5756291; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 246
TELECOMMUNICATION INFORMATION:
TELEPAN: 415-813-5600
TELEPAX: 415-44-0792
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TELEX: 706141
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATURE:
NAME/KEY: misc_difference
                                                                                                                                                                                                                                                                                                      181
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                  CITY: PALO ALTO STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
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                                                            US-08-484-192-16
                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY :
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LOCATION: replace(1, ")
OTHER INFORMATION: 'Anote- "This is a biotin-17
OTHER INFORMATION: nucleotide stretch of abasic residues."
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TITLE OF INVENTION: APPRAMERS SPECIFIC FOR BIOMOLECULES AND TITLE OF INVENTION: METHODS OF MAKING NUMBER OF SEQUENCES: 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
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US-09-016-520-35
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APPLICANT: DOWER, William J.
APPLICANT: CWIRLA, Steven A.
APPLICANT: CWIRLA, Steven A.
APPLICANT: UDFFIN, David J.
APPLICANT: WRIGHTON, Nicholas C.
APPLICANT: WAGSTROW, Christopher R.
TITLE OF INVENTION: PEPTIDES AND COMPOUNDS THAT BIND TO THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.0%; Score 0; DB 2; Length 2; 0.0%; Pred. No. 0; tive 0; Mismarches 1; Indel:
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ADDRESSEE: TOWNSEND AND TOWNSEND AND CREW STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 58
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND AND TOWNSEND AND CREW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : Two Embarcadero Center, 8th Floor
San Francisco
California
                                                                                                                                                                                                                                                                                                                                       16528A-024100
                                                                                                                                                                                                                                                                                            NAME: Garrett-Wackowsti, Eugenia
REGISTRATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 16528A-024
TELECOMUNICATION INFORMATION:
TELEPHONE: 415/576-0300
INFORMATION FOR SEQ 1D NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/726,464B
FILING DATE: 04-OCT-1996
CLASSIFICATION: 514
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                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           : Sequence 28, Application US/08726464B
; Patent No. 5932546
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COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: nucleic acid
                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 0.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 2 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                             COMPUTER READABLE FORM:
                                    San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                        California
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                                                                                            94111-3834
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                                                                          COUNTRY:
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                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                 Ouery Match 0.0%; Score 0; DB 2; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 37,330
REFERNCE/DOCKET NUMBER: 16528A-024100
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/576-0200
TELEPHONE: 415/576-0300
INFORMATION FOR SED ID NO: 28:
SEQUENCE CHARACTERISTICS:
NAME: Garrett-Wackowski, Eugenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 35, Application US/09016520A
; Patent No. 6127533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 35, Application US/09016520A; Patent No. 6127533; GENERAL INFORMATION:
                                                                                                                                                                                                       ; TOPOLOGY: linear
; MOLECULE TYPE: nucleic acid
US-08-726-464B-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                         LENGTH: 2 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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US-09-130-973-35
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LENGTH: 2
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APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Rawasaki, Andrew M
TITLE OF INVENTION: Aminoxy-Modified Oligonucleotides And Methods For
TITLE OF INVENTION: Aminoxy-Modified Oligonucleotides And Methods For
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (1)..(2)
OTHER INFORMATION: 2'-dimethylaminooxyethyl thymidine (T-2'-DMAOE)
OTHER INFORMATION: Description of Artificial Sequence: No. 6172209e1
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.0%; Score 0; DB 3; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                            ; LOCATION: (1)..(2); OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy (08-09-016-520-35)
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FILE REFERENCE: ISIS2824
CURRENT PAPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
BARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTWARE: PATENTIN VEY: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                 SEQ ID NO 35
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TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides And Methods For TITLE OF INVENTION: Making Same TITLE OF INVENTION: Making Same FILE REFERENCE: 1SIS295 CURRAW APPLICATION NUMBER: US/09/130,973 CURRAWY FILING DATE: 1998-08-07 NUMBER OF SEQ ID NOS: 20 Seg TONOS: 2.1
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APPLICANT: Mancharan, Muthiah
APPLICANT: Mancharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INFORTION: Anihooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS28A
CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
PRIOR PLING DATE: 1998-01-30
PRIOR FILING DATE: 1997-02-14
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! CCATION: (1)..(2)

OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy

US-09-477-092-35
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Matches 0; Conservative 0; Mismatches
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; Patent No. 6194598
APPLICANT: Mancharan, Muthiah
APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
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                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
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SOFTWARE: PatentIn Ver. 2.1
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US-09-477-902-35
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Search completed: August 24, 2002, 21:57:39 Job time: 21597 sec

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(without alignments)
56.110 Million cell updates/sec
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
                                                                     OM nucleic - nucleic search, using sw model
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. gb_gss:*
em_gss_hum:*
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8: em_htc:*

gb_est2:* gb_htc:* gb_est1:*

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Homo sapiens mRNA; EST DKFZp434J0410_s1 (from clone DKF2p434J0410)
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
               HSM003869 standard; RNA; EST; 1 BP.
                                                                                                                                                                                             EST; expressed sequence tag.
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Best Local Similarity 0.0%
Matches 0; Conservative
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Best Local Similarity (
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                           Duesterhoeft A., Lauber J., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                    Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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This clone is available at the RZPD in Berlin
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0; Mismatches
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Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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NH10B: sites NotI + SalI"
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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DH10B; sites Not1 + SalI"
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                                           Duesterhoeft A., Lauber J., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                 This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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                                                                      Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:9606"
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/clone_lib="434 (synonym: htes3). Vector pSportl; host
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens mRNA; EST DKFZp434N037_rl (from clone DKFZp434N037)
                                                                                                                                                                                                                                                                                                                  This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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                                                                                                                                                         Koehrer K., Beyer A., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                                              Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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                                                                                                                                                                                                                                                      Clone from S. Wiemann, sequenced by {\tt BMFZ} within the cDNA sequencing consortium of the German Genome Project No s1 sequence available
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12-WAR-1999 (Rel. 59, Last updated, Version 1)
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Matches 0; Conservative 0; Mismatches
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EST; expressed sequence tag.
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Butheria; Primates; Catarrhini; Hominidae; Homo.
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/clone="DKEZp4346017"
/clone_lib="434 (synonym: htes3). Vector pSport1; host
DH10B; sites NotI + SalI"
/dev_stage="adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERWANY
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Best Local Similarity 0.0%;
"...c 0; Conservative
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HSM011115
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Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /Clone_lib="586 (synonym: hutel). Vector pSportl; host
NH108; sites NotI + SalI/MluI"
/dev_stege="adult"
                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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                                    1; Indels
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       Score 0; DB 2; Length 1;
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
Query Match
0.0%; Scote v, 2-
Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
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Eutheria; Primates; Catarrhini; Homin:dae; Homo.
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/clone_lib="434 (synonym: htes3). Vector pSportl; host
DH10B: sites Not! + Sal!"
/dev_stage="adult"
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/clone=lib=#434 (synonym: htes3). Vector pSportl; host DH10B: sites NotI + SalI"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                0; Gaps
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email clone@rzpd.de
Please contact the RZPD: Ressourcenzeltrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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    /db_xref="taxon:9606"

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ID HSM011270 standard; RNA; EST; 1 BP.
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stage subtraction cona library (hasp)"

/dev_stage="Seedling roots of scots pine were infected for 6 days with H. annosum"

/note="Vector: pT-Adv; Site_1: EcoRI; The subtractive hybridization cDNA library was constructed from scots pine roots infected for 6-days with mycella of Heterobasidon annosum (FP5)."

1 others
                                                                                                                                                                                            stage subtraction cDNA library (hasp)"
/dev_stage="Seedling roots of scots pine were infected for
                                                                             /organism="Pinus sylvestris/Heterobasidion annosum"
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/clone="hasp002xj20f"
/clone="heterobasidion annosum - Scots pine infection
/clone_lib="Heterobasidion annosum - Scots pine infection
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/clone_lib="Heterobasidion annosum - Scots pine infection
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                                                                                                                                                                                                                                               6 days with H. annosum"

/note="Vector: pT-Adv; Site_1: EcoRI; The subtractive hybridization cDNA library was constructed from scots pine roots infected for 6-days with mycelia of Heterobasidon annosum (FP5)."

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Contact: Fred O. Aslegbu
Dept. of Forest Wycology & Pathology
Swedish University of Agriculture, Box 7026,S-750 07, Uppsala,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Pinus sylvestris/Heterobasidion annosum"
/db_xref="taxon:169015"
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100.0%; Pred. No. 0;
tive 0; Mismatches 0; Indels
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Eukaryota, mixed EST libraries.
1 (bases 1 to 1)
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Fax: +46 18 30 92 45
Email: Fred.Asiegbu@mykopat.slu.se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.0%; Score 0;
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Seg primer: T7 primer.
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hasp002x;20f Heterobasidion annosum - Scots pine infection stage subtraction cDNA library (hasp) Pinus sylvestris/Heterobasidion annosum cDNA clone hasp002x;20f, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/clone="DKF2p586F0B17"
/Clone_lib="586 (synonym: hutel). Vector pSportl; host
/Gl0Bs. sites NotI + SalI/MluI"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
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                                                                                   Homo sapiens mRNA; EST DKFZp586P0817_rl (from clone DKFZp586P0817)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                                                                                 Koehrer K., Beyer A., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases
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Pinus sylvestris/Heterobasidion annosum
Eukaryota; mixed EST libraries.
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Dept. of Forest Mycology & Pathology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: +46 18 67 15 98
Fax: +46 18 30 92 45
Email: Fred.Asiegbu@mykopat.slu.se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 bp
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B1817896 1 bp mRNA linear EST 04-OCT-2001 G3-023 Axolotl Lambda Zap Library Ambystoma mexicanum cDNA similar to Putative heterochromatin-like protein 1, mRNA sequence.
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Voss, S.R., King, D., Maness, N., Smith, J.J., Rondet, M., Bryant, S.V., Gardiner, D.M. and Parichy, D.M.
Expressed sequence tags from an axoloti limb regeneration library Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: srvoss@lamar.colostate.sdu
Single pass sequence from 5' end. Low quality sequence was trimmed
from the ends (PHRED error rate = 5%). Trace file available:
                                                                                                                                                                                                                                                                              Andystoma mexicanum
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibla; Batrachia; Caudata; Salamandroidea; Ambystomatidae;
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/orgalism="Ambystoma mexicanum"
/db_xref="taxon:8296"
/clone_lib="Axolot! Lambda Zap Library"
/tissue_type="Regenerating forelimb"
/dev_stage="Nedium-bud blastema"
0 a 1 c 0 g 0 t
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Location/Qualifiers
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Fort Collins, CO 80523, USA
Tel: 970 491 4869
Fax: 970 491 0649
                                                                                                                                                                                                                     BI817896.1 GI:15951508
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Department of Biology
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us-10-077-383-31.rge

Sequence 1 Sequence 2 Sequence 4 Sequence 5

Sequence

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Sequence 19 Sequence

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Minimum DB seq length: 0 Maximum DB seq length: 200000000

Searched:

US-10-077-383-31

Perfect score: Scoring table:

Run on:

Sequence:

Sequence 2 Sequence Primer. 6/1

Sequence Sequence

Sequence Sequence Sequence Sequence Sequence Sequence

Sequence DNA primer

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linear PAT 18-DEC-2001
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178656 Sequence 11
178657 Sequence 12
178658 Sequence 13
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AR026545 Sequence
AR026546 Sequence
AR026547 Sequence
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AX207477 Sequence
AR178168 Sequence
AR069073 Sequence
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AR026548 Sequence
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AR128994 Sequence
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100.0%;
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Ekins, R. Philip.
Unclassified.
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178.664 Million cell updates/sec
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                                   August 24, 2002, 21:54:45; Search time 2108.3 Seconds
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    GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                1797656 segs, 10463268293 residues
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Maximum Match 100%
Listing first 45 summaries
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gb_ba:* gb_htg:*

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Database :

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Query Score Match Length DB ID

Result

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linear PAT 29-MAR-1999
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DNA purification by triplex-affinity capture and affinity capture electrophoresis
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                                                          1 (bases 1 to 20)
Matson.R.S., Coassin,P.J., Rampal,J.B. and Caskey,C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5981185-A 93 09-NOV-1999;
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Reese, C.B. and Rao, M.V.
SULPHUR CONTAINING DINUCLEOTIDE PHOSPHORAMIDITES
PLEACH: WO 9729116-A 1 14-AUG-1997;
CRUACHEM LTD (GB)
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Sequence 1 from patent US 5482836.
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Arnold,L.J. Jr., Reynolds,M.A., Schwartz,D.A. and Daily,W.J.
Oligonucleoside cleavage compounds and therapies
Patent: US 5854410-A 23 29-DEC-1998;
Location/Qualifiers
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Matson, R.S., Coassin, P.J., Rampal, J.B. and Caskey, C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5881185-A 72 09-NoV-1999;
Location/Qualifiers
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Reese,C.B. and Rao,M.V.
BUDERHOR CONTAINING DINUCLEOTIDE PHOSPHORAMIDITES
Patent: WO 9729116-A 5 14-AUG-1997;
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Reses.C.B. and Rao,M.V.
LOLHUR CONTAINING DINUCLEOTIDE PHOSPHORAMIDITES
Patent: WO 9729116-A 4 14-AUG-1997;
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Reese, C.B. and Rao, M.V.
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Patent: WO 0110468-A 6 15-FEB-2001;
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/db_xref="taxon:32630"
/note="Synthetic Oligonucleotide"
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Krieg,A.M., Schetter,C. and Vollmer,J.C.
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Patent: WO 0122972-A 908 05-APR-2001;
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/db_xrsf="taxon:32630"
/note="Synthetic Oligonucleotide"
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THE GENERAL HOSPITAL CORPORATION (US)
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Sequence 5 from Patent W00110468.
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Gray, D.M. and Clark, C.L.
Method for selectively ranking sequences for antisense targeting
Patent: US 5856103-A 9 05-JAN-1999;
Location/Qualifiers
1 (bases 1 to 24)
Gray.b. M. and Clark.C.L.
Method for selectively ranking sequences for antisense targeting
Patent: US 5856103-A 8 05-JAN-1999;
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Maximum Match 100% Listing first 45 summaries Post-processing: Minimum Match 0%

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SIDSS/GGGdata/geneseq/geneseqn-embL/NA2001.DAT: 1: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1980.DAT:*
2: STDSS/gcgdata/geneseq/geneseqn-embl/Na1981.DAT:*
3: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1982.DAT:*
4: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1983.DAT:*
5: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1983.DAT:*
6: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1985.DAT:*
7: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1985.DAT:*
8: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1986.DAT:*
9: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1988.DAT:*
1/10: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1988.DAT:*
1/11: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1988.DAT:*
1/11: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1989.DAT:*
1/12: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1989.DAT:*
1/13: /SIDSS/gcgdata/geneseq/geneseqn-embl/Na1989.DAT:* N_Geneseq_032802:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	CAT gene target RN	W09526733 target s	Phosphorothicate o	Phosphorothicate o	Microsatellite seg	Immunostimulatory	Sequence of micros	Target sequence #2	Poly-pyrimidine ta
SUMMARIES	AAX91065	AAX56904	AAT86582	AAT86583	AAQ33557	AAF99703	AAQ33511	AAX00524	AAX00526
DB	2	16	18	18	13	22	13	20	20
a Query Match Length DB	18	20	21	21	22	22	23	24	24
& Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
Score	18	18	18	18	18	18	18	18	18
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0 00 0 00	11 12 13	18	100.0	24	20	AAX0052	7.2	Antisense Nucleic a	ense oligonuic acid trip
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00 0 00	7.7			7	22	AAF5799	7.	Nucleic	c acid tr
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υ υυ	16			24	22	AAF5800	0	Nucleic	c acid tr
υυ	17 18			24 24	22	AAF5800 AAF8778	1.1	Nucleic Hvbrid	ic acid tr d mRNA sec
υ	19			24	22	AAF8778	2	Hybr1d	DNA sedu
	21	180		24	22	AAF87784	34	Hybrid Hybrid	i mkna seque i DNA sequen
U C	22			26	17	AAT3277	& 0	Triple	helix-fc
ບ	24			26	20	AAZ3158	2.2	T7 PCR	primer.
o c	25			27	17	AAT0368	80 4	Triple	x-affinit
)	27			78	13	AAV6596	5.50	ol 1gor	Oligonucleotide t
	28		-	78	13	AAV1668	£ 5	01.190	ucleotide
	30			31	13	AA03352	. 0	Sequer	Sequence of micro
	31		-	31	13	AAQ3354	e :	Micros	Ę
	33			2 5	7 [AAO3355	c :-	Sequence	ice of micr
	34			33	13	AAQ3356	52.	Micros	Microsatellite se
ပ	35			36	12	AA01402	80 9	N-D-B	dc oligo
	37			2 2	7 -	AAQ3354	2.5	Micros	atellite
υ	38			80	22	AAH4687	. 72	B. nap	napus turgo
υ	39			39	20	AAX1887	0.	Maize	œ
υ	0.5			33	70	AAX1887	<u>_</u>	Maize	SSR oligonu
C	4.7			4 4	13	AAQ3361	ωu	Micros	Microsatellite
v	43			4	18	AAT4747	ŭ	Foldback	3 3
	44			41	15	AAQ7884	, m ¹	HCMV Ab	hea
						ALIGN	ALIGNMENTS		
RESULT	LT 1								
AAX91(1065/c AAX91065		standard;	RNA;	18 BI				
XX VX	AAX9106	ري							
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F Y	15-NOV-	-1999	(first	entry)	۲)				
DE I	CAT gene		target RNA		fragment				
×	940								
X X	binding af	aff	internucieosiayi inity; ss.	creos s.	1071	inkage;	; cnirality;	nybridization;	on; racemic;
SO VY	Syntheti	ic.							
X A	US5955597	97-A							
×									
G y	21-SEP-	-1999							
ΡF	30-JUN-1	1997	3076 ;	-088512	126.				
XX D	21 - NOV -	0	94118	-0343	5				
. K K ;	16-NOV- 30-JUN-	-1993	93US 93US 97US	-0154	013. 126.				
Y & S	(GENT-)	GENTA	TA INC.						
V I	Arnold	ĽJ,	Reynolds	s MA,	Ril	ley TA,	Schwartz DA	, Vaqhefi MM	<u>::</u>
XX DR	WPI; 19	99-5	39600/45						
XX									

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WO9729116-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT86582;
                                                                                                                                                                                                                                                             Query Match
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      8×888888888×8
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0
                                                                                               phosphonate internucleosidyl linkages of a preselected chirality which hybridize to a target RNA sequence. The method of making comprises: (a) synthesizing a nucleoside dimer, trimer, or tetramer with racemic internucleosidyl phosphonate linkages; (b) purifying the racemic nucleoside to a chirally pure nucleoside; and (c) sequentially linking at least 2 of the chirally pure nucleosides to form a synthetic oligomer that is enriched for phosphonate internucleosidyl linkages of a preselected chirality and is complementary to an RNA target sequence. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel oligonucleoside compound for hybridising to a target nucleic acid strand. The oligonucleoside comprises (a) an oligonucleoside sequence that is complementary to a target region or subregion of the target nucleic acid strand and (b) a portion that is non-complementary to a target site in the target region or subregion such that, when the oligonucleoside compound is hybridised to the target
                                                                                                                                                                                                                                                                       Rp chirally enriched synthetic oligomers have enhanced binding affinities for RNA compared to oligomers with racemic all methylphosphonate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cleavage; oligonucleoside; target; inter-strand orientation; inhibitor; disease; treatment; internucleotidyl bond cleavage; primer; ss.
                                                                                                                                                                                                                                                         methods are useful for providing chirally enriched synthetic oligomers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               strand, a base group at the site is oriented away from an inter-strand
                                                                                                                                                                                                                                                                                                             internucleosidyl linkagés. Sequences AAX31054-75 represent oligomers chemically synthesised using the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligo:nucleoside compounds for cleaving {\tt RNA} - having a sequence that is complementary to a target nucleic acid strand and a
                                                                         The invention provides methods for preparing oligomers having
                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 20; Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Daily WJ, Reynolds MA, Schwartz DA;
                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Scc...
100.0%; Pred. No. ...
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 0 A; 9 C; 0 G; 9 U; 0 other;
tetramers with enhanced binding affinities
                                        Example 19; Column 41-42; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 92; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9526733 target strand RNA 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX56904 standard; RNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18 AGAGAGAGAGAGAGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   non-complementary portion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1995-358439/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENT-) GENTA INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arnold LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX56904;
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orientation. The oligonucleoside and combinations are used for inhibiting
                               production of a selected protein in a cell by effecting cleavage at a site in a target region of cellular RNA that codes for the selected protein. They can be used for treating a condition in a mammal that is caused by the production of a selected protein. The oligonucleosides are target-mRNA specific and can be used against mRNA specific to a particular disease state. They are relatively harmless to non-targeted nucleic acid. The non-complementary unit enhances internucleotidyl bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a phosphorothioate oligonucleotide which was prepared by solid phase synthesis. The method comprises adding at least one dimeric phosphoramidite synthon, optionally having a protected thioester group in its internucleotide link, during the synthesis cycle. These novel dimeric phosphoramidite synthons are used as antisense molecules for inhibition of gene expression. The method gives increased yields of the phosphorothiate oligonucleotide (since fewer cycles are needed) and facilitates separation of impurities (greater difference in size compared with use of monomeric synthons).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Solid phase synthesis of phosphoro:thioate oligo:nucleotide(s) using new dimeric synthon(s) - useful as anti:sense molecules for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/note= "Phosphorothioate linkages between alternate
nucleotides (1 and 2, 3 and 4 etc.)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Phosphorothioate oligonucleotide; dimeric phosphoramidite synthon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            thioester; DNA synthesis; antisense oligonucleotide; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 18; DB 16; Length 20; 100.0%; Pred. No. 58; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 1 A; 9 C; 1 G; 9 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Phosphorothioate oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; Page 20; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT86582 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96GB-0002326.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 100.(
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CRUA-) CRUACHEM LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-415290/38.
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The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonuclectide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (Tf) n>9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp. microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic bovine DNA markers - used in genetic identification,
                                                                                                                                                                                                                                                                                                                          PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 18; DB 13; Length 22; 100.0%; Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        traits esp. in cattle, to allow selective breeding.
See also AAQ33501-34437.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 22 BP; 11 A; 0 C; 10 G; 0 U; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11 Similarity 100.0%; Pred. No. 58; 18; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                 Microsatellite sequence from clone AGLA248.
                                                                                                                                                                                                                                                                                                                                                         genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene mapping, and selective breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rable 7; Page 151; 517pp; English.
                                                                                                                                     AAQ33557 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF99703 standard; DNA; 22 BP.
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                  20 AGAGAGAGAGAGAGAG 3
Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1992-284684/34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9213102-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-AUG-1992.
                                                                                                                                                                                    AA033557;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a phosphorothioate oligonucleotide which was prepared by solid phase synthesis. The method comprises adding at least one dimeric phosphoramidite synthen, optionally having a protected thioester group in its internucleotide link, during the synthesis cycle. These novel dimeric phosphoramidite synthons are used as antisense molecules for inhibition of gene expression. The method gives increased yields of the phosphorothiate oligonucleotide (since fewer cycles are needed) and facilitates separation of impurities (greater difference in size compared with use of monomeric synthons).
                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOlid phase synthesis of phosphoro:thioate oligo:nucleotide(s) using new dimeric synthon(s) - useful as anti:sense molecules for inhibiting gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= a
/note= "Phosphorothioate linkages between alternate
nucleotides (1 and 2, 3 and 4 etc.)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phosphorothioate oligonucleotide; dimeric phosphoramidite synthon; thioester; DNA synthesis; antisense oligonucleotide; gene therapy;
                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                             Query Match 100.0%; Score 18; DB 18; Length 21; Best Local Similarity 100.0%; Pred. No. 58; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 18; DB 18; Length 21; 100.0%; Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
          Sequence 21 BP; 0 A; 10 C; 0 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 1 A; 10 C; 0 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Phosphorothioate oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 25; 38pp; English.
                                                                                                                                                                                                                                                                                                                                              AAT86583 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97WO-GB00327,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96GB-0002326.
                                                                                                                                                                               l agagagagagagagag 18
                                                                                                                                                                                                                                                                                                                                                                                                                                         ·25-MAR-1998 (first entry)
                                                                                                                                                                                                                        21 AGAGAGAGAGAGAGAG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CRUA-) CRUACHEM LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-415290/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reese CB;
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misc_difference 1

Synthetic.

AAT86583;

AAT86583/c

ò g 06-FEB-1997;

06-FEB-1996;

Rao MV,

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14-AUG-1997.

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0; Indels

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18;

Matches

Query Match

92WO-US00340. 91US-0642342.

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Georges M, Massey JM;
                                                                                                                                                                                                                                                                     WPI; 1992-284684/34.
                                                                                                                                                                                             (GENM-) GENMARK.
                                                                                                                                                        15-JAN-1991;
                                                                                                                   15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-OCT-1994;
                                         WO9213102-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5856103-A.
                                                                            06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JAN-1999
    Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX00524;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX00524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ŏ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulatery nucleic acid to a non-rodent subject in sufficient quantity to stimulater an nucleic acid. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects and/or orthomyxoviridae), bacterial antigens (e.g. horpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. horpesviridae, retroviridae staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a min to a Thi immune response and to activate immune cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                   Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vaccinating against tumors, infectious diseases, allergies and asthma
                                                                                                                                 infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a method for stimulating an immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Note: the present sequence may have a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 18; DB 22; Length 22; 100.0%; Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         using immunostimulatory Py-rich and TG nucleic acids -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence of microsatellite from clone AGLA209.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genetic mapping; traits; amplification; ss.
                                  Immunostimulatory nucleic acid #819.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Krieg AM, Schetter C, Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ33511 standard; DNA; 23 BP.
                                                                                                                                                                                                                                                                                     25-SEP-2000; 2000WO-US26383.
                                                                                                                                                                                                                                                                                                                                             27-SEP-1999; 99US-0156135.
23-AUG-2000; 2000US-0227436.
                                                                                                                                                                                                                                                                                                                             99US-0156113.
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12-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22 AGAGAGAGAGAGAGAG 5
                                                                                                                                                                                                                                                                                                                                                                                                                     COLE-) COLEY PHARM GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-273485/28.
                                                                                                                                                                                                           WO200122972-A2.
                                                                                                                                                                                                                                                                                                                          25-SEP-1999;
                                                                                                                                                                                                                                                05-APR-2001,
                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ33511;
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screening a genomic library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (Tf) >9 microsatellites in the bovine genome is estimated at >100. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequences upstream and downstream of the microsatellite sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                were used to generate the required PCR primers for in vitro amplification of the corresp. microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Target; antisense; selective rank; inhibition; ranking; stability;
                                                                                                                                                                                                 The sequence is that of a bovine microsatellite sequence obtd. by
Polymorphic bovine DNA markers - used in genetic identification,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 18; DB 13; Length 23; 100.0%; Pred. No. 58; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          traits esp. in cattle, to allow selective breeding.
See also AAQ33501-34437.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Target sequence #2 for antisense oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23 BP; 12 A; 0 C; 11 G; 0 U; 0 other;
                                      gene mapping, and selective breeding
                                                                                                                    Table 7; Page 132; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-0808474.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94US-0320507
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               interaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-MAR-1997;
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oligonucleotides (ASO) in a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises:

(a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA:ASO-DNA hybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid base pair types to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligoner binding interactions to select sequence zones for antisense targeting.
                                                                                                                                                                              This sequence represents a target mRNA for the generation of antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               efficiency comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and multiplying
efficiency - comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence cone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and multiplying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Target; antisense; selective rank; inhibition; ranking; stability;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 100.0%; Score 18; DB 20; Length 24; Local Similarity 100.0%; Pred. No. 58; es 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Selectively ranking nucleic acid molecules, for inhibitory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide for poly-purine target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 0 A; 12 C; 0 G; 12 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Column 13-14; 72pp; English.
                                                                                                                               Disclosure; Column 13-14; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX00525 standard; DNA; 24 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gray DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (TEXA ) UNIV TEXAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              interaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clark CL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX00525;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                              This sequence represents a target mRN. for the generation of antisense oligonucleotides (ASO) in a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises:

(a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA.ASO-UNA hybrid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of each of nearest-neighbour nucleic acid base pair types to provide of the nucleic acid base pair types to provide to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                     efficiency comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence some, substituting nearest-neighbour rucleic acid base pair fractions to determine the fractions and multiplying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Parget; antisense; selective rank; inhibition; ranking; stability;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Poly-pyrimidine target sequence for antisense oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 18; DB 20; Length 24; 100.0%; Pred. No. 58; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Selectively ranking nucleic acid molecules, for inhibitory
                                                                                                                                                                                 Selectively ranking nucleic acid molecules, for inhibitory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              select sequence zones for antisense targeting.
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                                                                                                                                                                                                                                                                                                                                          Disclosure; Column 13-14; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI: 1999-105098/09.
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                                                                                   Gray DM;
                              (TEXA ) UNIV TEXAS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-MAR-1997;
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                                                                                   Clark CL,
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targeted to a poly-purine mRNA sequence generated by a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises: (a) determining the fraction of each of a set of 3 nearest-neighbour nucleic acid base pair types in a target sequence of a nearest-neighbour nucleic acid base pair types in a target sequence of a nearest-neighbour nucleic acid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour of (c) multiplying the fractions of each of a series of 13 nearest-neighbour of a separate of 13 nearest-neighbour of (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense the process is used to rank nucleic acid sequences hased on the seminary and an anneary of the nucleic acid oligomer binding interactions to select
                                                                                                                                                                                                                                                                                                                                                                                                    o;
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Selectively ranking nucleic acid molecules, for inhibitory efficiency - comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and multiplying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotide for poly-pyrimidine target seguence.
                                                                                                                                                                                                                                                                                                                                                  Score 18; DB 20; Length 24;
Pred. No. 58;
                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                              Sequence 24 BP; 0 A; 12 C; 0 G; 12 T; 0 other;
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Matches 18; Conservative
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nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to select sequence zones for antisense targeting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA:DNA duplex capable of being used as an autisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Watson-Crick pairing. This is useful in antisense therapy, as it controls gene expression by causing transcription to be prevented. The present sequence is an example of a triplex sequence used in the methods of the invention.
                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNAasteriskbNa pyrimidineasteriskpurine duplex for use as an antisense molecule, by heating a triplex to dissociate a Watson-Crick paired
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antisense therapy; gene expression control; transcription; ss.
                                                                                                                                                                                  100.0%; Score 18; DB 20; Length 24;
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                                                                                                                                                                                                                       0; Indels
                                                                                                                              Sequence 24 BP; 12 A; 0 C; 12 G; 0 U; 0 other;
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                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                      AAF57996 standard; DNA; 24 BP.
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19-JAN-2000; 2000US-0487130.
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                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                               Local Similarity
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Gray DM, Hashem GM;
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              Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Generating nucleic acid molecule comprising Hoogsteen-paired
RNAasteriskDNA pyrimidineasteriskpurire duplex for use as an antisense
molecule, by heating a triplex to dissociate a Watson-Crick paired
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                                                                                                                                                                                                    Hoogsteen-paired duplex; Watson-Crick pairing; triplex; antisense therapy; gene expression control; transcription; ss.
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                                                                                                                                                                         Nucleic acid triplex DNA sequence #2.
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19-JAN-2000; 2000US-0487130.
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                                                                                   AAF57997 standard; DNA; 24
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24 AGAGAGAGAGAGAGAG
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Matches 18; Conserve
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                                                                                                                                                                                                                                                Synthetic.
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                                                        RESULT 13
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The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA:DNA duplex capable of being used as an antisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Matson Crick pairing. This is useful in antisense therapy, as it controls gene expression by causing transcription to be prevented. The present sequence is an example of a triplex sequence used in the methods of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an antisense molecule, by heating a triplex to dissociate a Watson-Crick paired
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antisense therapy; gene expression control; transcription; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 9; 23pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pyrimidine strand -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200105937-A2.
WO200105937-A2.
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The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA:DNA duplex capable of being used as an antisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Watson-Crick pairing. This is useful in transcription to be prevented. The present sequence is an example of a triplex sequence used in the methods of the invention.
                                        Generating nucleic acid molecule comprising Hoogsteen-paired
RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an antisense
molecule, by heating a triplex to dissociate a Watson-Crick paired
pyrimidine strand -
                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match
100.0%; Score 18; DB 22; Length 24;
Best Local Similarity 100.0%; Pred. No. 58;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                           Sequence 24 BP; 0 A; 12 C; 0 G; 12 U; 0 other;
                                                                                                                                               Claim 14; Page 17; 23pp; English.
  WPI; 2001-159523/16.
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Search completed: August 24, 2002, 22:08:17 Job time: 8850 sec

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0; Gaps

Appli Appl Appl

Sequence

35, A

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US-08-473-096-9 US-08-473-096-9 US-08-473-096-11 US-08-473-096-11 US-08-473-096-11 US-08-473-096-11 US-08-12-177A-385 US-09-387-300-35 US-09-387-300-10 US-09-285-309-10 US-08-285-309-10 US-08-285-309-10 US-08-379-556A-11 US-08-502-046-10 US-08-249-112-5

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Sequence 12, Appl
Sequence 4, Appli
Sequence 1, Appli
                                                                         (without alignments)
27.689 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23, Sequence 72, Sequence 93, Sequence 83, A
                                                            August 24, 2002, 21:57:39; Search time 159.68 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Description
                                                                                                                                                                                                                                                                                          GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-700-530-4
US-08-00-530-4
US-08-00-530-4
US-08-63-633-93
US-08-63-633-93
US-08-68-474A-9
US-08-68-474A-10
US-09-235-614-9
US-09-235-614-9
US-09-235-614-9
US-09-235-614-9
US-09-235-614-11
US-09-235-614-11
US-09-235-614-11
US-09-235-614-11
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US-09-235-614-11
US-09-235-614-11
US-09-235-615-11
US-08-267-8028-28
US-08-267-8038-46
US-08-267-8038-46
US-08-267-8038-46
US-08-473-096-1
US-08-473-096-1
                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
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                                                                                                                                                                            383533 segs, 122816752 residues
                                                                                                                                                                                                                                                                                                                                                                                                                     SUMMARIES
                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                         nucleic - nucleic search, using sw model
                                                                                                                                            IDENTITY_NUC Gapoo 10.0 , Gapext 1.0
                                                                                                                          1 agagagagagagag 18
                                                                                                  US-10-077-383-31
18
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Maximum DB seq length: 200000000
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Query
Match 1
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                                                                                                               Perfect score:
                                                                                                                          Sequence:
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RESULT 1 US-08-085-126-12/C Sequence 12, Application US/08085126A Patent No. 595557 GENERAL INFORMATION: APPLICANT: Riley, Timothy A. APPLICANT: APPLICANT: Anidy, Timothy A. APPLICANT: Control of the sequence of the se	Query Match 100.0%; Score 18; DB 2; Length 18; Best Local Similarity 100.0%; Pred. No. 5.2; Matches 18; Conservative 0; Mismatches 0; Indels 0; Gaps Qy 1 agagagagagagagag 18	RESULT 2 US-08-700-530-4/c ; Sequence 4, Application US/08700530 ; Patent No. 6316186 ; GENERAL INFORMATION:
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Sequence 3

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Sequence Seq

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Sequence Seq

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Gaps

0; Indels

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Pred. No. 5.2;
                                                Mismatches
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US-08-223-355-23/c
; Sequence 23, Application US/08223355
; Patent No. 5854410.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: other nucleic acid HYPOTHETICAL: Yes
                                                  ;
0
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RECISTRATION NUMBER: 32,274
REFERENCE/DOCKET NUMBER: 200,
TELECOMUNICATION INFORMATION:
TELEPHONE: 213/489-1600
TELEFAK: 213/955-0440
                         Best Local Similarity 100.0%;
Matches 18; Conservative 0
                                                                               l agagagagagagag 18
                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 611 W. Sixth St.
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 31-MAR-1994
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
                                                                                                    18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l agagagagagagagag 18
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100.0
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
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                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Cantor, Charles R. APPLICANT: Ito, Takashi APPLICANT: Smith, Cassandra L. TITLE OF INVENTION: DNA PURIFICATION BY TRIPLEX-AFFINITY TITLE OF INVENTION: CAPTURE AND AFFINITY CAPTURE ELECTROPHORESIS NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                        100.0%; Score 18; DB 4; Length 18; 100.0%; Pred. No. 5.2; tive 0; Mismatches 0; Indels
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                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Oligonucleotide 108-08-700-350-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DUCATION: 1..20
CTHER INFORMATION: /note= "BTC-20 Oligonucleotide, OTHER INFORMATION: biotinylated at 5'-end."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Karen S. Smith
STREET: 4 Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A-57666/KSS
PRIOR APPLICATION NUMBER: GB 9404709.9
PRIOR FILING DATE: 1994-03-11
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US/08/004,552
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/08004552
Patent No. 5482836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: A-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31,426
                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Smith, Karen S.
REGISTRATION NUMBER: 31,4
                                                        SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                      Ouery Match
Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: CA 94111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-004-552-1/c
                                                                       SEQ ID NO 4
                                                                                                              TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
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Gaps
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0
GENERAL INFORMATION:
APPLICANT: Arnold Jr., Lyle J.
APPLICANT: Reynolds, Mark A.
APPLICANT: Schwartz, David A.
APPLICANT: Daily, William J.
TITLE OF INVENTION: Oligonucleoside Cleavage Compounds and
TITLE OF INVENTION: Therapies
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 18; DB 2; Length 20; 100.0%; Pred. No. 5.2; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/223.355
                                                                                                                                                                                                                                                                         COUNTK1.

ZIP: 90017
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
"Counter: IBM PC Compatible
"Annew: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                        31-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; NAME/KEY: R183
; OTHER INFORMATION: target strand
US-08-223-355-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 72, Application US/08863639A
; Patent No. 5981185
; GENERAL INFORMATION:
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cuery Match 100.0%; Score 18; DB 2; Length 24; Best Local Similarity 100.0%; Pred. No. 5.2; Matches 18; Conservative 0; Mismatches 0. Trair
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APPLICATION UMBER: US/08/RDR 471*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 18; DB
Best Local Similarity 100.0%; Pred. No. 5.2
Matches 18; Conservative 0; Mismatches
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REGISTRATION NUMBER: 33,732
REFERENCE/DOCKET NUMBER: UTDAL:001
TELECOMNUNICATION INFORMATION:
TELEPHONE: (214) 740-6000
TELEFAX: (214) 740-8800
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : Locke Purnell Rain Harrell
2200 Ross Avenue, Suite 2200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 03-MAR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                            11859-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: Other nucleic acid US-08-863-639A-93
                                                                                                                                                     REFERENCE/OCCRET NUMBER: 11855
TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 796-6321
INFORMATION FOR SEQ ID NO: 93:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                         20,532
FILING DATE: May 28, 1997
CLASSIFICATION: 435
                                                ATTORNEY/AGENT INFORMATION:
NAME: Joseph E. Mueth
REGISTRATION NUMBER: 20,5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 agagagagagagagag 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
US-08-808-474A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            75201-6776
                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dallas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
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                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-808-474A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Diskette, 3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage COMPUTER: IBM compatible OPERATING SYSTEM: Windows 95 SOFTWARE: Corel WordPerfect 8 version CORRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Matson, Robert S.
APPLICANT: Coassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
APPLICANT: OLGGNUCLEOTIDE REPEAT ARRAYS
NUMBER OF SEQUENCES: 95
          APPLICANT: Matson, Robert S.
APPLICANT: Coassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
APPLICANT: Caskey, C. T.
NUMBER OF INVENTION: OLIGONUCLECTIDE REPEAT ARRAYS
NUMBER OF SEQUENCES: 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3: Sheldon & Mak
225 South Lake Avenue, 9th Floor
                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheldon & Mak
STREET: 225 South Lake Avenue, 9th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Corel WordPerfect 8 version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/863,639A FILING DATE: May 28, 1997 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11859-1
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; Sequence 93, Application US/08863639A
; Patent No. 5981185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; MOLECULE TYPE: Other nucleic acid US-08-863-639A-72
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hes 18; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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MEDIUM TYPE: Diskett
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MEDIUM TYPE: Diskette
                                                                                                                                                                          NUMBER OF SEQUENCES:
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| Patent No. 5856103
| GENERAL INFORMATION:
| APPLICANT: Gray, Donald M. APPLICANT: Clark, Chris L. TITLE OF INVENTION:
| TITLE OF INVENTION: FOR ANTISENSE TARGETING
100.0%; Score 18; DB 2; Length 20; 100.0%; Pred. No. 5.2;
                                       0; Indels
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23 AGAGAGAGAGAGAGAG 6
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                                                                                                                                                                                                                                                                                                                    24 base pairs
                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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TYPE: nucleic acid
STRANDEDNESS: single
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US-08-808-474A-10
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US-08-808-474A-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                      APPLICANT: Gray, Donald M.
APPLICANT: Glark, Chris L.
TITLE OF INVENTION: METHOD FOR SELECTIVELY RANKING SEQUENCES
NUMBER OF SEQUENCES: 37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 18; DB 2; Length 24; 100.0%; Pred. No. 5.2; tive 0; Mismatches 0; Indels
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COUNTRY: LEAGE
ZIP: 75201-6776
ZIP: 75201-6776
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBW PC COMPOTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHILIN Release #1.0, Version #1.30
SOFTWARE: PATCHILN Release #1.0, Version #1.30
SOFTWARE: OF MARR-1997
SOFTWARE: OF MARR-1997
ATTORNEY/AGENT INFORMATION:
NAME: Mayfield, Denise L.
REGISTRATION NUMBER: 33,732
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (214) 740-8800
INFORMATION FOR SEQ ID NO: 9:
CPONTENTE CHARRACTERISTICS:
                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Locke Purnell Rain Harrell
STREET: 2200 Ross Avenue, Suite 2200
                                                                               US-08-808-474A-9/c
; Sequence 9, Application US/08808474A
; Patent No. 5856103
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Sequence 10, Application US/08808474A
Patent No. 5856103
GENERAL INFORMATION:
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Matches 18; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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COMPUTER READABLE FORM:
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ZIP: 75201-6776
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US-08-808-474A-9
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Patent No. 5856103
GENERAL INFORMATION
APPLICANT: GTark, Chris L.
TITLE OF INVENTION: METHOD FOR SELECTIVELY RANKING SEQUENCES
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 18; DB 2; Length 24; 100.0%; Pred. No. 5.2;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION NUMBER: US/08/808,474A
FILING DATE: 03-MAR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/808,474A
FILING DATE: 03.MAR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                               NAME: Mayfield, Denise L.
REGISTRATION NUMBER: 33,732
REFERENCE/DOCKET NUMBER: UTDAL:001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (214) 740-8000
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Locke Purnell Rain Harrell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WIRESSEE: LOCKE PURUELL SUITE 2200 GTREET: 2200 Ross Avenue, Suite 2200
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OPERATING SYSTEM: PC-DOS/MS-DOS
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Best Local Similarity 100.0
Matches 18; Conservative
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APPLICANT: GRAY, DONALD M.
APPLICANT: CLARK, CHRISTOPHER L.
TITLE OF INVENTION: AN APPARATUS AND METHOD FOR SELECTIVELY RANKING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: GRAY, DONALD M.
APPLICANT: CLARK, CHRISTOPHER L.
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: Hybrid mRNA
US-09-235-614-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 18; DB 4; Length 24; 100.0%; Pred. No. 5.2; tive 0; Mismatches 0; Indels
  Best Local Similarity 100.0%; Pred. No. 5.2;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE OF INVENTION SEQUENCES FOR ANITSENED FILE REFERENCE: 91556/6684
CURRENT APPLICATION NUMBER: US/09/235,614
CURRENT FILING DATE: 1999-01-22
PRIOR APPLICATION NUMBER: 08/808,474
PRIOR FILING DATE: 1997-03-03
PRIOR FILING DATE: 1994-10-07
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATENTIN VEY. 2.1
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CURRENT APPLICATION NUMBER: US/09/235,614
CURRENT FILING DATE: 1999-01-22
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PRIOR FILING DATE: 1997-03-03
PRIOR APPLICATION NUMBER: 08/320,507
PRIOR FILING DATE: 1994-10-07
                                                                                                                                                                                                                   RESULT 13
US-09-235-614-10/c
Sequence 10, Application US/09235614
; Patent No. 6183966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 11, Application US/09235614 ; Patent No. 6183966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
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SOFTWARE: PatentIn Ver. 2.1
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Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                    24 AGAGAGAGAGAGAGAG 7
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LENGTH: 24
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APPLICANT: GRAY, DONALD M.
APPLICANT: CLARK, CHRISTOPHER L.
TITLE OF INVENTION: AN APPARATUS AND METHOD FOR SELECTIVELY RANKING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
TITLE OF INVENTION: SEQUENCES FOR ANTISENSE TARGETING
FILE REFERENCE: 91556/66184
CURRENT RILING DATE: 1999-01-22
PRIOR APPLICATION NUMBER: 08/808,474
PRIOR FILING DATE: 1997-03-03
PRIOR FILING DATE: 1998-01-07
                                                           0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: Hybrid mRNA US-09-225-614-8
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Query Match 100.0%; Score 18: DB 2; Length 24; Best Local Similarity 100.0%; Pred. No. 5.2; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 18; DB 4; Length 24; 100.0%; Pred. No. 5.2; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08/320,507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 9, Application US/09235614; Patent No. 6183966
                                                                                                                                                                                                                                                                                         ; Sequence 8, Application US/09235614; Patent No. 6183966
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: GRAY, DONALD M. APPLICANT: CLARK, CHRISTOPHER L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
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PRIOR FILING DATE: 1994-10-07
NUMBER OF SEQ ID NOS: 38
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                                                                                                                                         1 AGAGAGAGAGAGAGAG 18
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                              ; GENERAL INFORMATION:
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                                                                                                                                                                                                                                         RESULT 11
US-09-235-614-8
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LENGTH: 24
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Best Local Similarity

100.0%; Score 18; DB 4; Length 24;

Query Match

Query Match

0; Gaps

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ö;
                Gaps
                                                                                                                                                                             APPLICANT: CROUZET, JOEL
APPLICANT: CROUZET, JOEL
APPLICANT: SCHERRAIN, Daniel
APPLICANT: SCHERRAIN, Daniel
APPLICANT: WILS, Pierre
TITLE OF INVENTION: PURIFICATION OF A TRIPLE HELIX FORMATION
TITLE OF INVENTION: WITH AN IMMOBILIZED OLIGONUCLEOTIDE
OUNDER OF SEQUENCES: 25
CORRESSONDENCE ADDRESS:
ADDRESSED: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, Mailstop 3C43
CITY: Collegeville
                0;
           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match 100.0%; Score 18; DB 4; Length 26; Best Local Similarity 100.0%; Pred. No. 5.2; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION DATA:
APPLICATION NUMBER: US/08/860,038
        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDENESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION NAME
APPLICATION NUMBER: FR 94/15162
FILING DATE: 16-DEC-1994
PRIOR APPLICATION DATE: WE FR95/01468
FILING DATE: 08-NOV-1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 15
US-08-860-038-19/c; Sequence 19, Application US/08860038; Patent No. 6287762; GENERAL INFORMATION:
                               l agagagagagagagag 18
                                               18; Conservative
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 Matches
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Search completed: August 24, 2002, 21:57:39 Job time: 21597 sec

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August 24, 2002, 21:19:25; Search time 4810.92 Seconds (without alignments) 50.499 Million cell updates/sec
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                          13736207 segs, 6748477542 residues
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                                                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                 OM nucleic - nucleic search, using sw model
                                                                                                                                                                 US-10-077-383-31
18
1 agagagagagagagag 18
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em_gss_inv:*
em_gss_pln:*
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Maximum DB seq length: 2000000000
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2: em_esthum:*
3: em_estin:*
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em_estov:*
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em_estro:*
em_htc:*
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gb_htc:*
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Perfect score:
Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

em_gss_vrt:*

SUMMARIES

	Description	86 AZ489586 1M0322C1	52 AZ510952 1M0355G1	14 AZ983014 2M0264H02	66 AZ433566 1M0219C02	97 AZ492997 1M0327K2	57 AZ770557 1M0572N08	97 AZ394897 1M0158H1	98 AZ589098 1M0397B1	78 AZ627978 1M0476L04	54 AZ464354 1M0273M1	51 AZ633751 1M048911	.88 AZ430288 1M021401	.88 AZ822888 2M0096G06	26 AZ973926 2M0248N1	18 AZ389918 1M0151E05	24 AZ510124 1M0354K22	95 AZ659095 1M0536E18
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	Query Match Length DB	19	19	19	20	20	20	21	21	21	22	22	23	23	23	25	52	25
æ	Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
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AZ564804 1M0545H24 AZ342914 1M0076C22 AZ675959 1M0367010 AZ66145 1M0540500 AZ666145 1M05408C00 AZ666145 1M05408C00 AZ771239 1M057815 AZ771239 1M057815 AZ8338913 AZ338913 AZ6338913 1M049082 AZ6338913 AZ01090820 AZ65531 1M049082 AZ65531 1M049082 AZ66043 AZ028117 AZ65946 1M0271N04 AZ641732 1M050117 AZ65946 1M0278117 AZ65946 1M0278117 AZ661735 1M0504123 AZ61789 1M0504123 AZ61781 1M0604123 AZ61732 1M0504123 AZ61732 1M0504123 AZ61732 1M0504123 AZ616033 1M0428114 AZ63463 1M0438114 AZ63463 1M043812 AZ61613 1M043812 AZ61613 1M043812 AZ61613 1M043812 AZ61613 1M043812 AZ61613 1M043172 AZ61613 1M043172 AZ61613 1M043172	linear GSS 05-OCT-2000 cy Mus musculus genomic ctebrata; Euteleostomi; muridae; Murinae; Mus. Duval,B., Hamil,C., E., Pedersen,T., Reilly c, von Niederhausern,A. end reads from 10kb ., 20 S. 2030 E., SLC, UT
AZ564804 AZ342914 AZ541486 AZ65145 AZ671439 AZ71239 AZ71239 AZ71239 AZ732991 AZ632991 AZ622991 AZ622991 AZ622991 AZ65531 AZ65531 AZ65531 AZ65531 AZ65531 AZ65531 AZ65531 AZ656035 AZ86035 AZ66035 AZ656035 AZ666035	DNA CLM libran Ce. urognathl. "Meenen," Tingey,A th paired arch Bldg
	ALIGNMENT 2F Mouse 10kb plasmid UUG GCIM0322C12 F, DNA sequen 1 GI:10659481 uuse. uulus a: Metazoa; Chordata; Cra i: Lot 19) set 1co 19; set 1co 10; set 2co 1co 10; set 1
33333331110099888877779999999999999999999999999	S86 2012F Mouse 10kb 2010C1M032C12 F 586.1 GI:106594 mouse. usculbrain; Rameria; Robert B. We right, D., Welse, R. Whole genome scillahed (2000) ct: Robert B. We reity of Utah 08, Blomedical P. We Sandray 08, Blomedical P. We Sandray 1058 5606 801 585 7177 1058 5606 801 585 7177 11 19 /organism="Market Corfurgray 1 198
	AZ489586 AZ4805 AZ489586 AZ489586 AZ489586 AZ489586 AZ489586 AZ489586 AZ489586 AZ489586 AZ480586 AZ4806 AZ4806 AZ4806 AZ4806 AZ4806 AZ4806 AZ4806 AZ4806 AZ48
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0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	RESULT 1 AAA 89586/C LOCUS DEFINITION ACCESSION VERSION VERSION VERSION VERSION TEPTERENCE ONCANISM TITLE JOURNAL COMMENT COMMENT SOUICE

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/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                      Laboratory Mouse DNA Resource
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Insert Length: 10000 Std Erro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            δy
                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polywucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                                                                                           electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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                                                                                                                                                                                                                                                                                                                                                                                                                             adaptored vector DNA, and transformed into chemically-competent E. coli XLi0-Gold (Stratagene) cells and selected for ampicillin resistance. ^{\circ}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ510952 19 bp DNA linear GSS 05-0CT-20C 1M035015N Mouse 10kb plasmid UUGCIM library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                  /lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 18; DB 12; Length 19; 100.0%; Pred. No. 3.5e+03;
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                                                                                             Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ddunn@genetics.utah.edu
Length: 10000 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. nc.
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Seq primer: CACAGGAAACAGCTATGACC
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University of Utah Genome Center
University of Utah
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Location/Qualifiers
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/clone="UUGC1M0355G15"
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/sex="Male"
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Fax: 801 585 7177
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                                                                                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, M., Stokes, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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/lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-"/hote="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
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/organism="Mus musculus"
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ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
                                                                      /lab_host="E. coli strain XL10-Gold, T1-resistant, F-"
/hote="Vector: PWD42Dry, Purified genomic DNA from M.
musculus C57BL/6J (female) was obtained from the Jackson
Laboratory Mouse DNA Resource
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Musmalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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                      /clone_lib-"Mouse 10kb plasmid UUGC2M library"
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University of Utah Genome Center
University of Utah
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/clone="UUGC2M0264H02"
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                                                                                                                   /lab_host="E. Coli strain XLIO-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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Seg primer: CGTTGTAAAACGACGCCAGT
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University of Utah Genome Center
University of Utah
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/db_xref="taxon:10090"
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                               /clone="UUGC1M0219C02
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Plate: 0327 row: K c
                                                                                       /sex="Male"
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Matches 18; Conservative
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Inter://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.065 inch orifice at constant velocity. The sheared DNA was blunt end repeated with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwW42 (gil473114 [giPAT29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A2394897 21 bp DNA linear GSS 03-0CT-200 IM0158H11R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0158H11 R, DNA sequence.
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                                                                                                                                                                                                                                        musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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                                                                                                                                                                                                              /note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                    /clone_lib="Mouse 10kb plasmid UUGC1M library"
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100.0%; Pred. No. 3.5e+03;
tive 0; Mismatches 0;
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Insert Length: 10000 Std Error: 0.00
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   /organism="Mus musculus"
                                                          /db_xref="taxon:10090"
/clone="UUGC1M0572N08"
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                                /strain="C57BL/6J
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AZ394897.1 GI:10509969
                                                                                                                                              /sex="Male"
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Fax: 801 585 7177
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,M., Stokes,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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1M0572N08F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0572N08 F, DNA sequence.
                                                                                                                                           /lab_host="E. Coli strain XLIO-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                       /clone_lib="Mouse 10kb plasmid UUGClM library"
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Insert Length: 10000 Std Error: 0.00
Plate: 0572 row: N column: 08
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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High quality sequence stop: 20.
Location/Qualifiers
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared bNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar axcess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwbs. (qi[4732114]qb]ARI29072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chamically-competent E. coll XL10-Gold (Stratagene) cells and selected from mpicillin resistance."
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Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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1M0476L04F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
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                                                                                                                                                                                                                             /lab_host="E. Coli strain XLIO-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                          /clone_lib="Mouse 10kb plasmid UUGC1M library"
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100.0%; Pred. No. 3.5e+03;
                                                                                                                                                                                                                                                                                                                              Laboratory Mouse DNA Resource
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lnsert Length: 10000 Std Error:
Plate: 0476 row: L column: 04
Seg primer: CGTTGTAAAACGACGGCCAGF
Class: plasmid ends
                                                      /organism="Mus musculus"
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University of Utah
                                                                                                             /db_xref="taxon:10090"
/clone="UUGC1M0397B19"
Location/Qualifiers
                                                                             /strain="C57BL/6J
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Contact: Robert B. Weiss
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  electrophoresis. Vector DNA was prepared from a derivative of pwD42 (q1|4732114|qb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                             /lab_host="E. Coli strain XLIO-Gold, T1-resistant, F-"
hote="Vector: PRN2ADY; Purified genomic DNA from M.
musculus G57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Fesource
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Insert Length: 10000 Std Errcr: 0.00
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                      /ordanism="Mus musculus"
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University of Utah Genome Center
University of Utah
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/clone="UUGC1M0158H11"
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                                                   /strain="C57BL/6J
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DEFINITION

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1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                        /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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Insert Length: 10000 Std Error: 0.00
Plate: 0273 row: M column: 11
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Seq primer: CACACAGGAAACAGCTATGACC
                                                                                /organism="Mus musculus"
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University of Utah Genome Center
                                                                                                                                  /db_xref="taxon:10090"
/clone="UUGC1M0476L04"
High quality sequence stop: 21.
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Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a was hydrodynamically sheared by repeated passage through a was blunt end repeated with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWM42 (gil473114 [gb]AR129072.1], a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coll KLI0-Gold (Stratagene) cells and selected for ampicilin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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/sex="Male"
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Best Local Similarity 100.0%; Pred. No. 3.5e+03;
Matches 18; Conservative 0; Mismatches 0; Indels
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University of Utah Genome Center
University of Utah
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                                                                                                                                        /strain="C57BL/63
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Fax: 801 585 7177
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Plate: 0214 row: O column: 12
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Tel: 801 585 5606
Fax: 801 585 7177
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,M., Rose,M., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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1M0214012R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0214012 R, DNA sequence.
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  Seg primer: CGTTGTAAAACGACGGCCAGT
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                     High quality sequence stop: 22.
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                                                                            Location/Qualifiers
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Fax: 801 585 7177
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http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 lnch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pw042 (gil4732114|gb|AF129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance.
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University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Musinae; Lico 23)
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2M0096G06R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC2M0096G06 R, DNA sequence.
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                                                                                                                     /organism="Mus musculus"
High quality sequence stop: 23.
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/clone="UUGC1M0214012"
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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2M0248N15F Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0248N15 F, DNA sequence.
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Seq primer: CACACAGGAAACAGCTATGACC
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Contact: Robert B. Weiss
University of Utah Genome Center
                                                                                                     High quality sequence stop: 23.
Location/Qualifiers
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/clone="UUGC2M0096G06"
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Insert Length: 10000
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/note="vector: PWD42Nv; Purified genomic DNA from M. musculus C57BL/G1 (female) was obtained from the Jackson Laboratory Mouse DNA Resources (documents/dnares/). The DNA was hydrodynamically sheared by repeated gassage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymerated to the blunt ends in high molar excess. The adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0248 row: N column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                            /organism="Mus musculus"/strain="C57BL/6J"
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/clone="UUGC2M0248N15"
                                                                                                                                                                                   High quality sequence stop: 23. Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /sex="Female"
                                                                                                                                                    Class: plasmid ends
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charotactory mouse two measources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.065 inch orlifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gli4732114[g5]ARI29072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-"
/lote="Vector: PWD42nv Puiffied genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/db.xref="taxon:10000"
/clone="UUGCLMO151E05"
/clone=11b="Mouse 10kb plasmid UUGCLM library"
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100.0%; Score 18; DB 12; Length 25;
Best Local Similarity 100.0%; Pred. No. 3.5e+03;
Matches 18; Conservative 0; Mismatches 0; Indels
                           Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0151 row: E column: 05
Seq primer: CGTGTAAAACGACGCCAGT
Class: plasmid ends
High quality sequence stop: 25.
Location/Qualifiers
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Perfect score:

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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Neisserial antigens
Patent: WO 9924578-A 25 20-MAY-1999;
PIZZA MARIAGRAIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRADIG GUIDO (IT); MASIGNANI VEGA (IT)
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Sequence 25 from Patent WO9924578.
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/db_xref="taxon:32644"
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178.664 Million cell updates/sec
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                                                                                  August 24, 2002, 21:54:45; Search time 2108.3 Seconds
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          GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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PAT 07-SEP-2000

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Neisserial antigens
Patent: w0 924578-A 35 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Plaza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Neisserial antigens
Patent: MO 9924578-A 45 20-MAY-1999;
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Sequence 35 from Patent WO9924578.
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Sequence 45 from Patent WO9924578.
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DEFINITION Sequence 45 from Patent WO9924578.
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Location/Qualifiers
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Neisserial antigens
Patent: WO 9924578-A 35 20-MaY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
Location/Qualifiers
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Patent: WO 9924578-A 25 20-MAY-1999;
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Sequence 25 from Patent WO9924578.
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Paten: WO 924578-A 45 20-MAY-1999;
Pizza MARIAGRAZIA (IT): SCARLATO VINCENZO (IT): RAPPUOLI RINO (IT):
CHIRON SPA (IT): GRANDI GUIDO (IT): MASIGNANI VEGA (IT)
Location/Qualifiers
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Patent: WO 9924578-A 55 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Sequence 55 from Patent WO9924578.
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza Harigens
Pizza Mariagazia (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GIUDO (IT); MASIGNANI VEGA (IT)
Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V. Neisserial antigens
Patent: WO 9924578-A 55 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Neisserial antigens
Patent: WO 9924578-A 73 20-MAY-1999;
PIZZA MARIAGRAIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT): GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
Location/Quanifiers
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Pizza,M., Scarlato,V., Rappuoli,R., Grandi,G. and Masignani,V.
Nelsserial artigens
Patont: WO 9924578-A 109 20-MAY-1999;
PATONT: WO 9924578-A 109 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Patent: WO 9924578-A 109 20-MAY-1999;
PIZZA MARIAGRAZIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Neisserial antigens
Patent: WO 9924578-A 91 20-MAY-1999;
PIZZA MARIAGRAIA (IT); SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT);
CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pizza, M., Scarlato, V., Rappuoli, R., Grandi, G. and Masignani, V.
Pizza Mariagharia, Pizza Mariagharia, Pizza Mariagharia, Rip), SCARLATO VINCENZO (IT); RAPPUOLI RINO (IT); CHIRON SPA (IT); GRANDI GUIDO (IT); MASIGNANI VEGA (IT)
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

/SIDS5/gcgdata/genesec/geneseqn-embl/NA2001A.DAT:*/SIDS5/gcgdata/geneseq/geneseqn-embl/NA2001B.DAT:*/SIDS5/gcgdata/geneseq/geneseqn-embl/NA2002.DAT:* /SIDS5/gcgdata/geneseq/geneseqn-embl/NA2000.DAT:*

Description	Human mutant KCNO3	Human mutant KCNO3	Sequence encoding	Sequence encoding	Plasmid pEZZ318.th	Plasmid pEZZ318.th	CAPL trinucleotide	CAPL trinucleotide	Human TUB gene pro
SUMMARIES	AAX57131	AAX57131	AAN80743	AAN80743	AAQ85491	AAQ85491	AAT33326	AAT33326	AAA94655
DB DB	20	20	6	6	16	16	17	17	21
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                                                                            This invention describes novel human and mouse potassium channel proteins KCNQ2 and KCNQ3. Detecting mutations in sequences that encode KCNQ2 and KCNQ3. Detecting mutations in sequences that encode KCNQ2 or the loss of one copy of these genes, is used for diagnosis and prognosis of benign familial neonatal epilepsy (REWE). Calsonial myotonic epilepsy (JME) or rolandic epilepsy (RE). Cells (or transgenic animals) that express wild-type or mutanit KCNQ2 or 3 (also the proteins themselves in cell-free form) are used to screen for agents that can be used to treat or prevent these forms of epilepsy. Fragments of the encoding mutations or for isolation of related sequences, while the complete sequences may be used in gene therapy to provide wild-type protein Antibodies specific for mutant or wild-type proteins are used as diagnostic reagents and for drug screening. The KCNQ2 and 3 proteins are useful in rational design of drugs and therapetically (in proteins
                                                                                                                                                                                                                                                                                                                                                                                          replacement theraples). The forms of epilepsy associated with mutations in KCNQ2 and 3 sequences can now be diagnosed early (before symptoms are manifest), and better treatment options will be available.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KCNQ2; KCNQ3; human; murine; potassium channel; diagnosis; prognosis; benign familial neonatal epilepsy; BRNE; juvenile myotonic epilepsy; JME; rolandic epilepsy; mutant; treatment; screening; epilepsy; detection; gene therapy; drug screening; primer; ss.
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Nucleic acid encoding potassium channels KCNQ2 and 3
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                                          Claim 65; Page 151; 195pp; English.
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CC KCNQ2 or KCNQ3, or the loss of one copy of these genes, is used for diagnosis and prognosis of benign familial neonatal epilepsy (BFNB).

CC diagnosis and prognosis of benign familial neonatal epilepsy (BFNB).

CC transgenic animals) that express wild-type or mutant KCNQ2 or 3 (also the proteins themselves in cell-free form) are used to screen for agents that can be used to treat or prevent these forms of epilepsy. Fragments of the encoding nucleic acids are used as probes or primers, either for encoding mutations or for isolation of related sequences, while the complete sequences may be used in gene therapy to provide wild-type protein. Antibodies specific for mutant or wild-type proteins are used as diagnostic reagents and for drug screening. The KCNQ2 and 3 proteins are useful in rational design of drugs and therapeutically (in replacement therapies). The forms of epilepsy associated with mutations mainfest, and better treatment options will be available.

AAX57074-X57139 are primers used in the method of the invention.
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                                    A DNA segment with a nucleotide sequence from about 130 to about 918 of the sequence given here is claimed. Also claimed are antibodies which immunoraect with hurfi and the claimed peptides (given in AAP80713). The antibodies may be used in immunoassays for detection of hurfi. The claimed peptides may be used to inhibit the binding of hurf to coagulation factor VII/VIIa in vivo.
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Disclosure; ; pp; English.
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Matches 1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mat_peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein inhibitors of serine proteinase(s), e.g. furin, deriv. from turkey ovomucoid third domain - used as laboratory reagents to study the proteinase(s), or as chemotherapeutic agents to treat diseases associated with them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OMTKV3; turkey ovomucoid third domain peptide inhibitor; furin inhibitor; protease inhibitor; pEZZ318.thy2; probe; Escherichia coli; ss.
                                                                         0; Indels
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          Length 3;
   0.0%; Score 0; DB 9;
100.0%; Pred. No. 0;
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Query Match 0.0%; Score 0; DB 9
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                          AAQ85491 standard; cDNA; 3 BP
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96WO-US02108

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16-FEB-1996;
                          17-FEB-1995;
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                                                                                                         Von Hofe E;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         In synthetic analogs of turkey ovomucoid domain protein (6-56) (OMTKY3), given in AAR69818-24, the region immediately adjacent to the reactive site peptide bond is mutated to include the consensus sequence of furin or other serine protease. Polynucleotides encoding such analogs are incorporated into pEZZ318.thy2 and expressed in Escherichia coli RV308 (ATCC 31608).
                                                                                                                                                                                                                                                                                                                                                                              Protein inhibitors of serine proteinase(s), e.g. furin, deriv. from turkey ovomucoid third domain - used as laboratory reagents to study the proteinase(s), or as chemotherapeutic agents to treat diseases associated with them
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                                                                                         OMTKY3; turkey ovomucoid third domain peptide inhibitor;
furin inhibitor; protease inhibitor; pEZ2318.thy2; probe;
Escherichia coli; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
0.0%; Score 0; DB 16; Length 3;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 42-45; 66pp; English.
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                                      18-AUG-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    osteosarcoma; therapy; ss.
                                                                                                                                                                                                                                                                                                                            Anderson S, Laskowski M;
                                                                  Plasmid pEZZ318.thy2.
                                                                                                                                                                                                                                                                                                  (LASK/) LASKOWSKI M.
                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-066900/09.
                                                                                                                                                                                                                                                                                      (ANDE/) ANDERSON S.
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                                                                                                                                                                                                                                                          09-JUL-1993;
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              AAQ85491;
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                                                                                                                                                                                  Agrawal S, Engebraaten O, Fodstad O, Hovig E, Maelandsmo GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Agrawal S, Engebraaten O, Fodstad O, Hovig E, Maelandsmo GJ;
                                                                                                                                                                                                                                                                                                                                                                 Synthetic oligo:nucleotide(s) inhibiting CAPL gene expression -
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                                                                                                                                                                                                                                                                                                                                                                                                         useful to inhibit metastatic cancer, partic. osteosarcoma
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Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
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                                                                                                       (NORA-) NORWEGIAN RADIUM HOSPITAL RES FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2; Page 56; 70pp; English.
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95US-0391375.
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                                                                           (HYBR-) HYBRIDON INC
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Sequence 3 BP; 0 A; 0 C; 3 G; 0 U; 0 other;

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                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
Novel antisense oligonucleotides capable of inhibiting CAPL gene expression may include the trinucleotide GOC (ART33326, given in 5' to 3' direction) found in codon 14 of CAPL mRNB. These and other antisense oligonucleotides (AAT3337-36) complementary to specific regions of the CAPL gene (see also AAT3335), as well as CAPL-specific ribozynes (AAT33337-40) can be administered to a patient as a means of inhibiting metastatic cancer.
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                                                                                                                                                                                                                                                                                                                    unery Match 0.0%; Score 0; DB 17; Length 3; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                      Sequence 3 BP; 0 A; 1 C; 1 G; 1 U; 0 cther;
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neurosensory defect associated gene family. The novel cDNAs are mouse tub form I (see AAA94629), mouse tub form II (see AAA94630), human TUB form I (see AAA94630), human TUB form I (see AAA94630), human TUB form I (see AAA94630), human TUB I (see AAA94635), human TULPI (see AAA94635), human TULPI (see AAA94636), human TULPI (see AAA94638). The novel coding sequences are useful as immunogens to raise antibodies that specifically identify TUBFYTULP defects. The novel proteins encoded by the present sequence can be used for the treatment of neurosensory defects. The novel proteins encoded by the present sequence can be used dystrophies. The present sequence is a probe used to isolate the novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalian proteins expressed in retina and brain, useful for producing antibodies and for diagnosing neurosensory defects including cochlear degeneration, peripheral retinal degeneration and cone-rod retinal
                                                                                                                                                                                                      Human; TULP; neurosensory defect; retina; retinal dystrophy; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to human and murine cDNAs from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       North M, Nishina P, Noben-Trauth K, Naggert J;
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                                                                            AAA94655 standard; DNA; 3 BP.
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                                                                                                                                            (first entry)
                                                                                                                                                                         Human TUB gene probe #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AXYS-) AXYS PHARM INC.
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                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                          15-JAN-2001
                                                                                                                                                                                                                                                                                                                                                27-FEB-1998;
                                                                                                                                                                                                                                                                                   US6114502-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to human breast cancer expressed polynucleotides ARMO07849-ARMISOR99) and methods of assessing whether a patient is afflicted with breast cancer be examining the correlation between the expression of certain markers and the cancerous state of breast cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                Human breast cancer expressed polynucleotide 12701.
                                                                                                                                                                                                                                            Human; breast cancer; cell marker; cytostatic; ss.
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tive 0; Mismatches
0.0%; Pred. No. 0; ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lillie J, Xu Y, Wang Y, Steinmann K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 2245; 3695pp; English.
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                                                                                                                                   AAL20244 standard; cDNA; 3 BP.
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29-MAR-2000; 2000US-0193480.
15-MAY-2000; 2000US-0193480.
09-JUN-2000; 2000US-021315.
25-JUL-2000; 2000US-0220534.
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              Conservative
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 Best Local Similarity
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                                                                                                                                                                                                                                                                       Homo sapiens
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The invention relates to human breast cancer expressed polynucleotides (AAL07544-AAL26789) and methods of assessing whether a patient is sefflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells. The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic
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                                                                    Human breast cancer expressed polynucleotide 12701.
                                                                                                                                        Human; breast cancer; cell marker; cytostatic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lillie J, Xu Y, Wang Y, Steinmann K;
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15-MAY-2000; 2000US-0205230.
09-JUN-2000; 2000US-0211315.
25-JUL-2000; 2000US-020534.
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07-DEC-2001 (first entry)
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                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                         New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
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0.0%; Pred. No. 0;
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                                                                 94WO-US06884.
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                                                                                                                                                                           (PHAR-) PHARMAGENICS INC.
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07-JAN-1994;
                                                              17-JUN-1994;
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07-JAN-1994;
                     05-JAN-1995.
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The sequences given in AAQ81642-95 are oligonucleotides which modulate to activity of basic fibroblast growth factor (bFGF) by binding to the bFGF protein. Most easy. The sequences given in AAQ81664-67 represent core sequences of which at least one is present in each of the binding oligos. These oligonucleotides may form a single strand, double strand, a stem-loop structure, a bubble structure, a pseudoknot or a closed, circular structure. bFGF binds to high affinity receptor and low affinity heparin-like molecules on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell surface. These oligonucleotides bind to bFGF in competition with its receptor and heparin. These oligonucleotides may inhibit or enhance the activity of bFGF. Particularly, they inhibit necowascularisation so they can be used to suppress growth of solid tumours and to reduce the risk of metastasis. They can be used as diagnostic reagents to determine the presence of thrombin, or used in
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                                                                                                                                                                                                                   New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
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(PHAR-) PHARMAGENICS INC.
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                                                                  Beutel BA, Joesten ME;
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The sequences given in AAQ81642-95 are oligonucleotides which modulate the activity of basic fibroblast growth factor (bFGF) by binding to the bEGF protein. Most esp. the sequences given in AAQ81664-67 represent core sequences of which at least one is present in each of the binding oligos. These oligonucleotides may form a single the binding oligos. These oligonucleotides may form a single stead, double strand, a stem-loop structure, a bubble structure, a pseudoknot or a closed, circular structure. BFGF binds to high a finity receptor and low affinity heparin-like molecules on the affinity receptor and heparin. These oligonucleotides may inhibit or chance the activity of bFGF. Particularly, they inhibit or enhance the activity of bFGF. Particularly, they inhibit or evascularisation so they can be used to suppress growth of solid tumours and to reduce the risk of metastasis. They can be used as chippersents of thrombin, or used in
New oligo-nucleotide(s) that bind to basic fibroblast growth factor - modulating, esp. inhibiting, its activity, useful in treating cancer, preventing metastasis, and diagnosis.
                                                                                                                    Claim 3; Page 25; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy.
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Query Match 0.0%; Score 0; DB 16; Length 4; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels Search completed: August 24, 2002, 22:08:17 Job time: 8850 sec 1 n 1 1 c 1 οy οp

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1; Indels

Sequence 4 BP; 0 A; 2 C; 1 G; 1 U; 0 other;

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27.689 Million cell updates/sec
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Sequence 702, App
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                                                                                                                                                                       August 24, 2002, 21:57:39; Search time 159.68 Seconds
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/cgn2_6/ptcdata/2/ina/%B_COMB.seq:*
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/cgn2_6/ptcdata/2/ina/%B_COMB.seq:*
/cgn2_6/ptcdata/2/ina/%B_COMB.seq:*
/cgn2_6/ptcdata/2/ina/ma/ma/kfiles1.seq:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-US93-(0977-702
US-08-268-679B-8
US-08-268-679B-8
US-08-457-274A-16
US-08-441-192-16
US-08-441-192-16
US-08-464-28
US-08-726-464B-28
US-08-726-464B-28
US-09-130-973-35
US-09-130-973-35
US-09-105-335
US-09-477-902-35
US-09-477-902-35
US-09-411-862A-15
US-09-411-862A-15
US-09-411-862A-15
US-09-411-862A-15
US-09-411-862A-16
US-09-791-213D-46
US-07-791-213D-46
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US-07-791-213D-46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            383533 seqs, 122816752 residues
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                                                                                                              OM nucleic - nucleic search, using sw model
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                                                                                                                                                                                                                                                                                        US-10-077-383-33
18
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Maximum DB seq length: 200000000
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Sequence:
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Sequence Sequence

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                                                           Sequence 2, A Sequence 2, A Sequence 46, Sequence 46,
                                                                                                                                                                                                                                                                                       Sequence 702, Application PC/TUS9300977
GENERAL INFORMATION:
TITLE OF INVERTION:
NUMBER OF SEQUENCES: 711
                         Sequence 7
Sequence 2
Sequence 2
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Sequence 2
Sequence 2
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                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Knobbe, Martens, Olson, and Bear
STREET: 620 Newport Center Dr. Sixteenth Floor
CITY: Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PAPLICATION NUMBER: PCT/US93/00977
FILING DATE: 19930129
US-07-791-213D-62

US-08-266-679B-7

US-08-265-743-22

US-08-295-743-22

US-08-602-036A-2

US-08-602-036A-2

US-08-602-036A-2

US-08-293-150A-46

US-08-293-150A-62

US-08-293-150A-62

US-08-293-150A-62

US-08-293-150A-62

US-08-293-150A-62

US-08-293-150A-62

US-08-293-150A-62

US-08-642-407A-2

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US-08-642-407A-2

US-08-642-407A-2

US-08-642-407A-2

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US-08-643-709-9

US-08-03-365A-36
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Pred. No. 0;
0; Mismatches
                                                                                                                                                                                                                                  ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Altman, Daniel E.
REGISTRATION NUMBER: 34,115
REFERENCE/DOCKET NUMBER: HITACHI.006H
TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 702: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: CDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.08;
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Best Local Similarity 0.0%;
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MEDIUM TYPE: Floppy disk
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EDNESS: double
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ZIP: 92660
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PCT-US93-00977-702
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PCT-US93-00977-702
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us-10-077-383-33.rni

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APPLICATION NUMBER: US/08/268,679B
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POSITION IN GENOME: N.A.
US-08-268-679B-8
                                                                                                                                                                           MARIA C.H. LIN
                                                                                                                                                                                                                                                                                                                                                                                         SINGLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: FLOPPY
                                                                                                                                                                                                                                                                                                                                                                    NUCLEIC ACID
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MOLECULE TYPE:
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                                     Sequence 702, Application PC/TUS9300977
GENERAL INFORMATION:
TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
NUMBER OF SEQUENCES: 711
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
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                                                                                                                                   SEE: Knobbe, Martens, Olson, and Bear
: 620 Newport Center Dr. Sixteenth Floor
Newport Beach
                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IRP PC compatible
COMPUTER: IRP PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/00977
FILING DATE: 19930129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: WIMMER, ECKARD; MOLLA,
APPLICANT: AKHTERUZZAMAN; PAUL, ANIKO V.
TITLE OF INVENTION: DE NOVO CELL-FREE
TITLE OF INVENTION: SYNTHESIS PICORNAVIRUS
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: HITACHI.006H
TELECOMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
TELEFRAX: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P. STREET: 345 PARK AVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORD PERFECT # 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 8, Application US/08268679B
; Patent No. 5674729
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 702:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E.
REGISTRATION NUMBER: 34,1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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NEW YORK
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CLASSIFICATION:
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    RESULT 2
PCT-US93-00977-702/c
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PCT-US93-00977-702
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                                                                                                                                                                                                                                     92660
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                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                  STATE: CA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-268-679B-8
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APPLICANT: AKHTENGZAMAN; PAUL, ANIKO V.
TITLE OF INVENTION: DE NOVO CELL-FREE
TITLE OF INVENTION: SYNTHESIS PICORNAVIRUS
CORRESCEDIES: GORGES: GORGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0887-4095 US2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: MORGAN & FINNEGAN, L.L.P.
345 PARK AVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPENAL:
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/08/268,679B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PC-DOS/MS-DOS
FILING DATE: 30-UNN-1994
PRIOR APPLICATION DATA:
RAPLICATION NUMBER: 07\846,914
FILING DATE: 06-M8R-1992
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07\719,761
FILING DATE: 24-JUN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8, Application US/08268679B Patent No. 5674729
                                                                                                                                                                                                                                                                           07\\719,761
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: FLOPPY DISK
IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 29,323
REPERENCE/COCKET NUMBER: 08
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (212) 758-4800
TELERAX: (212) 751-6849
TELEX: 421792
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 07\71 FILING DATE: 24-JUN-1991 ATTORNEY/AGENT INFORMATION:
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STRAIN: Learn-PyR
DEVELOPMENTAL STAGE: Adult
POSITION IN GENOME:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14603
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                                                                  ; CHROMOSOME,
US-08-457-274A-16
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                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Scott, Jeffrey G.
APPLICANT: Tomita, Takashi
TITLE OF INVENTION: Cytochrome P450lfr Gene and Its Uses
CORRESPONDENCE ADDRESS: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.0%; Score 0; DB 1; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/457,274A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19603/240 (D-1519)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Nixon, Hargrave, Devans & Doyle
STREET: P.O. Box 1051, Clinton Square
CITY: Rochester
REGISTRATION NUMBER: 29,323
REFERENCE/DOCKET NUMBER: 0887-4095 US2
TELECOMMULICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELERAX: (212) 751-6849
TELERAX: 42179
INFORMATION FOR SEQ IN 0: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 16, Application US/08457274A
; Patent No. 5734086
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY AGENT INFORMATION:
NAME: Goldman, Michael L.
REGISTRATION NUMBER: 30,727
REFERENCE/DOCKET NUMBER: 1960:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 716-263-1304
                                                                                                                                                                                                                                            DESCRIPTION: OLIGONUCLEOTIDE POTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Musca domestica
                                                                                                                                                                                                                                                                                                ORIGINAL SOURCE: N.A.
POSITION IN GENOME: N.A.
US-08-268-679B-8
                                                                                                                                                                                            SINGLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
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                                                                                                                                                                      NUCLEIC ACID
                                                                                                                                                                                                        TOPOLOGY: UNKNOWN
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                                                                                                                                                                                      STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Scott, Jeffrey G.
APPLICANT: Tomita, Takashi
TITLE OF INVENTION: Cytochrome P4501pr Gene and Its Uses
WUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                 Ouery Match 0.0%; Score 0; DB 1; Length 2; Best Local Similarity 0.0%; Pred. No. 0; Matches 0; Conservative 0; Mismatches 1; Indel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CNERBY APPLICATION DATA:
APPLICATION NUMBER: US/08/457,274A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30,727
IER: 19603/240 (D-1519)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Nixon, Hargrave, Devans & Doyle
STREET: P.O. Box 1051, Clinton Square
CITY: Rochester
CHROMOSOME/SEGMENT: Chromosome 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN: Learn-PyR
DEVELORMENTAL STRAGE: Adult
POSITION IN GENOME:
CHROMOSOME/SEGMENT: Chromosome 1
                                                                                                                                                                                                                                                                                                                                                      US-08-457-274A-16/c
; Sequence 16, Application US/08457274A
; Patent No. 5734086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 716-263-1304
TELEPAX: 716-263-1600
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Musca domestica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Goldman, Michael L. REGISTRATION NUMBER: 30,7 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                H: 2 base pairs
nucleic acid
DEDNESS: double
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us-10-077-383-33.rni

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LOCATION: replace(1, "")

CTHER INFORMATION: /note= "This is a biotin-17

OTHER INFORMATION: nucleotide stretch of abasic residues."
US-08-484-192-16
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                                                                                    TITLE OF INVENTION: APTAMERS SPECIFIC FOR BIOMOLECULES AND TITLE OF INVENTION: METHODS OF MAKING NUMBER OF SEQUENCES: 181
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

0.0%; Score 0; DB 1; Length 2;
Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches 1; Indels
Matches 0; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  246102002221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/934,387
FILING DATE: 21-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: BARRETT, Ronald W.
APPLICANT: CWIRLA, Steven A.
APPLICANT: CWIRLA, Steven A.
APPLICANT: JOHNSON, Sherril S.
APPLICANT: WIGHTON, Nicholas C.
APPLICANT: DUFFIN, David J.
                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 28, Application US/08726464B Patent No. 5932546
                                                                                                                                                                                ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-813-5600
TELEFAX: 415-494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: GRACEY, NANCY J. REGISTRATION NUMBER: 28,216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS: LENGTH: 2 base pairs TYPE: nucleic acid
LATHAM, JOHN
LEUNG, LAWRENCE
                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                          VERMAAS, ERIC
TOOLE, JOHN J
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CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                             94304
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                 APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                           COUNTRY:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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CTHER INFORMATION: /note= "This is a biotin-17
CTHER INFORMATION: nucleotide stretch of abasic residues."
US-08-484-192-16
                                                                                                                                                                                  APPLICANT: ALBRECHT, GLENN
APPLICANT: LATHAN, JOHN
APPLICANT: LATHAN, JOHN
APPLICANT: VERMAAS, ERIC
APPLICANT: TOOLE, JOHN J
TITLE OF INVENTION: APTAMERS SPECIFIC FOR BIOMOLECULES AND
TITLE OF INVENTION: METHODS OF MAKING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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PRICR APPLICATION DATA:
APPLICATION NUMBER: US 07/934,387
FILING DATE: 21-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/484,192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-484-192-16/c
; Sequence 16, Application US/08484192
; Patent No. 5756291
                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: MORRISON & FOERSTER STREET: 755 PAGE MILL ROAD
                                                                                         Sequence 16, Application US/08484192; Patent No. 5756291
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TELECOMMUNICATION INFORMATION: TELEPHONE: 415-813-5600
TELEPAX: 415-44-0792
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APPLICANT: GRIFFIN, LINDA C.
APPLICANT: ALBRECHT, GLENN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 706141
INFORMATION FOR SEQ ID NO: 16:
                                                                                                                                 GENERAL INFORMATION:
APPLICANT: GRIFFIN, LINDA C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94304
                                                                     US-08-484-192-16
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                          ; MOLECULE TYPE
US-08-726-464B-28
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APPLICANT: CWIRLA, Steven A.
APPLICANT: WISHIA, Steven A.
APPLICANT: JOHNSON, Sherril S.
APPLICANT: WISHIGHON, Nicholas C.
APPLICANT: WAGSTROM, Christopher R.
APPLICANT: WAGSTROM, Christopher R.
TITLE OF INVENTION: THROMBOPOIETIN RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
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                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/726,464B
FILING DATE: 04-0CT-1996
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PAPLICATION NUMBER: US/08/726,464B
FILING DATE: 04-OCT-1996
ADDRESSEE: TOWNSEND AND TOWNSEND AND CREW STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                               ATTORNEY AGENT INFORMATION:
NAME: Garrett-Wackowski, Eugenia
REGISTRATION NUMBER: 37 330
REFERENCE/POCKET NUMBER: 16528A-024100
TELECOMMUNICATION INFORMATION:
TELECHOME: 415/576-0200
TELEFAX: 415/576-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
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ADDRESSEE: TOWNSEND AND TOWNSEND AND CREW
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Matches 0; Conservative 0; Mismatches
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IBM PC compatible
iYSTEM: PC-DOS/MS-DOS
                                                                                                                                   Eloppy disk
IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: nucleic acid
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            STREET: TWO Entroca
CITY: San Francisco
CTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid
EDNESS: single
                                                                                            ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
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California
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CLASSIFICATION:
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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APPLICANT: Cook, Phillip D
APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
ITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (1)..(2)
CTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
US-09-016-520-35
NAME: Garrett-Wackowski, Eugenia
REGISTATION NUMBER: 37,330
REFERENCE/DOCKET NUMBER: 16528A-024100
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/576-0200
TELEPRAX: 415/576-0300
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Patent No. 6127533
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                                                                                                                                                           INFORMATION FOR SEQ ID NO: 28: SEQUENCE CHARACTERISTICS: LENGTH: 2 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                   single
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us-10-077-383-33.rni

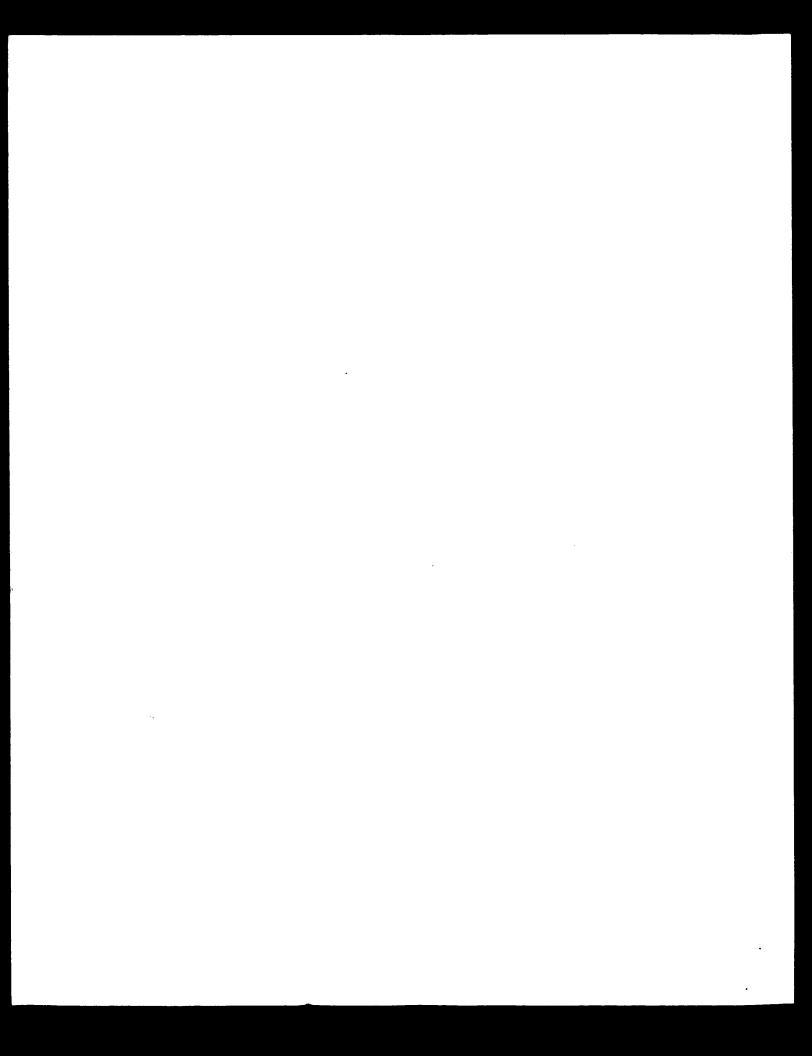
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FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn Ver. 2.1
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PRIOR APPLICATION WUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
PRIOR APPLICATION NUMBER: 60/037,143
PRIOR FILING DATE: 1997-02-14
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; Patent No. 6194598
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APPLICANT: Manoharan, Muthiah
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NUMBER OF SEQ ID NOS: 47
SOFTWARE: PatentIn Ver. 2.1
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APPLICANT: Cook, Phillip Dan
APPLICANT: Cook, Phillip Dan
APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Rawasaki, Thazha P
APPLICANT: Rawasaki, Andrew M
TITLE OF INVENTION: Making Same
TITLE OF INVENTION: Making Same
FILE REFERENCE: ISISASAS
CURRENT FILING DATE: 1989-09-07
NUMBER OF SEO ID NOS: 28
SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
SOFTWARE: PATENTING VOS: 47
SOFTWARE: PATENTIN VOS: 2.1
SEQ ID NO 35
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FILE REFERENCE: ISIS2824
                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
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APPLICANT: COCK Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Rawashi, Andrew P
APPLICANT: Rawashi, Andrew P
TITLE OF INVENTION: Aninoxy-Modified Oligonucleotides And Methods For
TITLE OF INVENTION: Aninoxy-Making Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (1)..(2)
OTHER INFORMATION: 2'-dimethylaminooxyethyl thymidine (T-2'-DMAOE)
OTHER INFORMATION: Description of Artificial Sequence: No. 6172209e1
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APPLICANT: Manoharan, Muthiah
APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2834
CURRENT APPLICATION UNDER: US/09/477,902
CURRENT APPLICATION NUMBER: 09/016,520
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Search completed: August 24, 2002, 21:57:39 Job time: 21597 sec



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50.499 Million cell updates/sec
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Copyright (c) 1993 - 200) Compugen Ltd.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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X1-2blue; sites Noti + SalI"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                         0;
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                                                                                                                                                                                                                                                                                                                                                                                                                          Clone from S. Wiemann, sequenced by GBF within the cDNA sequencing consortium of the German Genome Project
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12-WAR-1999 (Rel. 59, Last updated, Version 1)
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X1-2blue; sites NotI + SalI"
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Enkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
12-MAR-1999 (Rel. 59, Created)
12-MAR-1999 (Rel. 59, Last updated, Version 1)
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12-MAR-1999 (Rel. 59, Created)
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Matches 0; Conservative 0; Mismatches
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DH10B; sites NotI + SalI"
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                                                              Duesterhoeft A., Lauber J., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                    Clone from S. Wiemann, sequenced by Qiagen within the cDNA sequencing consortium of the German Genome Project also available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059 Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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This clone is available at the RZPD ir Berlin
Please contact the RZPD: Ressourcenzertrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERWANY; Email: clone@rzpd.de
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostom1; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone_lib="434 (synonym: htes3). Vector pSport1; host
DH10B; sites Not1 + Sal1"
/dev_stage="adult"
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/dev_stage="adult"
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                       Enkaryota; Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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DH10B; sites NotI + SalI"
/dev_stage="adult"
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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/clone="DKF2p434G017"
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                                       Homo sapiens (human)
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1; Mammalia;
Butheria; Primates; Catarrhin1; Hominidae; Homo.
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hlubs; sites NotI + SalI"
/dev_stage="adult"
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This clone is available at the RZPD in Berlin
Please contact the RZPD: Ressourcementrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Matches 0; Conservative 0; Mismatches
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MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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Matches 0; Conservative 0; Mismatches
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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h1108; sites NotI + SalI"
/dev_srage="adult"
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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12-WAR-1999 (Rel. 59, Last updated, Version 1)
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Matches 0; Conservative
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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DH10B; aites Not! + SalI/Mlu!"
Gaps
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Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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/clone_"hasp002xj20f"
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                                                                       /organism="Pinus sylvestris/Heterobasidion annosum'
                                                                                                                                                                      stage subtraction cDNA library (hasp)"
                                                                                                 /db_xref="taxon:169015"
                       Location/Qualifiers
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Location/Qualifiers
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  Seq primer: T7 primer
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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DH10B; sites NotI + SalI/MluI"
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Dept. of Forest Mycology & Pathology
Swedish University of Agriculture, Box 7026,S-750 07, Uppsala,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                     Homo sapiens mRNA; EST DKF2p586P0817_rl (from clone DKF2p586P0817)
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Please contact the R2PD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
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                                                                                                                                                                                                                                                                                                                                                             Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
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12-MAR-1999 (Rel. 59, Created)
12-MAR-1999 (Rel. 59, Last updated, Version 1)
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Best Local Similarity 0.0%; Pred. No. 0;
Matches 0; Conservative 0; Mismatches
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Fax: +46 18 30 92 45
Email: Fred.Asiegbu@mykopat.slu.se
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/dey_stage="Seedling roots of scots pine were infected for
/dey_stage="Seedling roots of scots pine were infected for
/dey_station cDNA library was constructed from scots
pine roots infected for 6-days with mycelia of
pine roots infected for 6-days with mycelia of
Heterobasidon annosum (PF5)."

let no constructed for 6-days of the constructed from scots
/dey_station cDNA library was c
/dev_stage="Seedling roots of scots pine were infected for 6 days with H. annosum //nete="Vector: pT-Adv; Site_1: EcoRI; The subtractive hybridization cDNA library was constructed from scots pine roots infected for 6-days with mycelia of
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Dept. of Forest Mycology & Pathology
Swedish University of Agriculture, Box 7026,S-750 07, Uppsala,
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Pinus sylvestris/Heterobasidion annosum
Eukaryota: mixed EST libraries.
1 (bases 1 to 1)
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Best Local Similarity 100.0%; Pred. No. 0;
1. Conservative 0; Mismatches
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Best Local Similarity 100.0%; Pred. No. u;
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Email: Fred.Asiegbu@mykopat.slu.se
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BI817896 1 bp mRNA linear EST 04-OCT-2001 G3-O23 Axolotl Lambda Zap Library Ambystoma mexicanum cDNA similar to Putative heterochromatin-like protein 1, mRNA sequence.
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(bases 1 to 1)

Voss,S.R., King,D., Maness,N., Smith,J.J., Rondet,M., Bryant,S.V.,
Gardiner,D.M. and Parichy,D.M.

Expressed sequence tags from at axolot1 limb regeneration library
Unpublished (2001)
Contact: Voss S.R.

Department of Biology
Colorado State University
Fort Collins. CO 80523, USA
Tel: 970 491 4869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: srvoss@lamar.colostate.edu Single pass sequence from 5' erd. Low quality sequence was trimmed from the ends (PHRED error rate = 5\%). Trace file avallable:
                                                                                                                                                                                                                                                                                                              Ambystoma mexicanum
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Caudata; Salamandroidea; Ambystomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
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/tissue_type="Regenerating forelimb"
/dev_stage="Medium-buc blastema"
1 c 0 g 0
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Location/Qualifiers
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